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415.756 Million cell updates/sec
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Listing first 45 summaries
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Pred. No. 18 No. is the number of results predicted by chance to have a greater than or equal to the score of the result being printed, serived by analysis of the total score distribution.

SUMMARIES

24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	σı	σ	4	ω	N	_	No.	Regn 1+
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Abu66618 Human PRO	Human	Abo17664 Novel hum	Abb82287 Human LP8	Abp52427 Human LP8		Aea28787 Human int	Aea50049 Full leng	Adx58772 Interleuk	Adw64518 Human IL-		Ads18363 Human Zcy	Human	Adq88067 Human Zcy	Adm95041 Human Int		Human	Abw00889 Human cyt			Abg67190 Inflammat		Aab85259 Human int	Aay08661 Human Zcy	- 1 1 1	

45	44	43	42	41	40	9	38	37	36	ω 5	34	33	32	31	30	29	28	27	26	77
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Adb14586	Ada91523	Adb16431	Ada87229	Ada79090	Ada96786	Ada85574	Adb30278	Ada67271	Ada47476	Adb15690	Ada86126	Adb27647	Adb19106	Ada61321	Ada18698	Ada76048	Ada45617	Abu66894	Abo24889	ADUSY699
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ALIGNMENTS

RESULT 1 AAY08661 Zcyto10; four alpha helix cytokine; vulnerary; cytostatic; haemostatic; anti-inflammatory; anti-asthmatic; growth factor; maintenance factor; trachea; salivary gland; stomach; panoreas; muscle; gene therapy; cancer; proliferation; differentiation; modulator; gastrointestinal tract; oral cavity; asthma; tracheobronchial tract; bronchitis; wound healing; platelet count; thrombocytopenia; human. AAY08661; 26-NOV-1997; 25-NOV-1998; 03-JUN-1999. WO9927103-A1 Homo sapiens Human Zcyto10 allele 1 protein. 09-AUG-1999 AAY08661 standard; protein; 176 (ZYMO) ZYMOGENETICS INC. (first entry) 97US-00979156 98WO-US025228 Ā

New mammalian cytokine-like polynucleotide useful in the treatment of asthma and bronchitis.

Conklin DC,

Haldeman

BA,

Grossmann

۸,

1999-357840/30. DB; AAX77692.

Claim 1; Page 64-65; 82pp; English.

This invention describes a novel mammalian four alpha helix cytokine designated Zcytol0 which has vulnerary, cytostatic, anti-inflammatory, anti-asthmatic and haemostatic activity. Zcytol0 may be a growth and/or maintenance factor in the trachea and salivary glands, stomach, pancreas and muscle. Zcytol0 polynucleotides are useful as sources of primers and probes, and to determine if the Zcytol0 gene is present on chromosome 1, or to detect any mutations that may have occurred. Zcytol0 polynucleotides are also useful as gene therapy reagents. The Zcytol0 and

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CC identified as having binding affinity to Zcytol0, should be useful in the
CC treatment of conditions associated with abnormal physiology or
CC development, including abnormal proliferation, e.g. cancerous conditions,
CC useful for modulating cell proliferation, cell differentiation or
CC cytokine production in the prevention or treatment of conditions
CC characterized by abnormal cell proliferation, cell differentiation or
CC cytokine production. Zcytol0 polypeptides, agonists and antagonists may
CC characterized by seeful in the regeneration of the gastrointestinal
CC tract or oral cavity, as well as in the treatment of asthma, and other
CC diseases of the tracheobronchial tract such as bronchitis. Zcytol0
CC polypeptides may also be useful to promote wound healing, and to increase
CC platelet count, e.g. in cancer patients who experience thrombocytopenia
CC due to chemotherapy or radiation therapy
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                                                                                                  Isolated subunits treating
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antiinflammatory; antipsoriatic; antiasthmatic; antibacterial;
dermatological; antiulcer; antagonist.
                                                                                                                                                                                                                                               WPI; 2001-398320/42.
                                                                                                                                                                                                                                                                                                               Foster DC,
Rixon MW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-DEC-1999; 99US-00471774
22-JUN-2000; 2000US-0213416P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human interleukin-20 (IL-20) receptor polypeptide
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                                      Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                     (ZYMO ) ZYMOGENETICS INC.
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                                                                                                         inflammatory diseases
                                                                                                                                         IL-20RA and
                                                                                                                                                                                                                                                                                                               Presnell
                                                                                                                                                                                                                                                                                                                                           Xu W,
                                   Page 53-54; 119pp; English
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100.0%; Pred. No. 5e-88;
:ive 0; Mismatches
                                                                                                                                     20 soluble receptor comprising two polypeptide IL-20RB, useful for down-regulating IL-20 and thus
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                                                                                                                                                                                                                                                                                                                  BA;
                                                                                                         such as psoriasis
                                                                                                                                                                                                                                                                                                                                              Kelly JD,
                                                                                                                                                                                                                                                                                                                                              Sprecher CA,
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Best Local S
                                                                                                                                                                                                                       Thompson
Sprecher
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; interleukin-20; II-20; antagonist; psoriasis; eczema; de adult respiratory disease; asthma; bronchitis; pneumonia; multiple organ failure; inflammatory lung injury; septic shock; bacterial pneumonia; inflammatory bowel disease; rheumatoid art ulcerative colitis; Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammatory diseases such as psoriasis, inflammatory lung injury such asthma or bronchitis, adult respiratory disease (ARD), septic shock, multiple organ failure, bacterial pneumonia, eczema, atopic and contact dermatitis, and inflammatory bowel disease such as ulcerative colitis as Crohn's disease. Sequences AAB85259-262 represent human IL-20 receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            comprising two polypeptide subunits IL-20RA (formerly known as ZcytoR7) and IL-20RB (formerly known as DIRS1). The two subunits are preferably linked together. In one embodiment, one subunit is fused to the constant region of the light chain of an immunoglobulin, and the other subunit is fused to constant region of the heavy chain of an immunoglobulin. The light chain and the heavy chain are connected via a disulphide bond. The soluble receptor can be used to down-regulate IL-20 and thus treat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human interleukin-20, IL-20, protein #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAU04049;
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                                                                                                                                                                                                                                                                                                                                                              23-DEC-1999; 99US-00470898.
22-JUN-2000; 2000US-0213341P.
                                                                                                                                                                                                                                                                                                                                                                                                                                               22-DEC-2000; 2000WO-US035305
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                                                                                                                                                                                                                                                                                           (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   121
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Blumberg
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100.0%; Pred. No. 5e-88;
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                                                                                                                                                                                                                                                 Wenfeng
                                                                                                                                                                                                                 Eagan
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Jaspers S
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                                                                                                                                                                                                                                                       Kelly JD;
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Treating interleukin-20 induced inflammation in a mammal, such as adult respiratory disease, eczema, psoriasis, contact dermatitis, multiple organ failure and septic shock, involves administering IL-20 antagonist.

WPI; 2001-418045/44

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Best Local S
Matches 176
                                                                                                                                                                                                                                                                                                              Inflammation; interleukin-20; IL-20; interleukin-8; IL-8; chemokine; neutrophil; monocyte; basophil; eosinophil; chemoattractant; psoriasis; periodontal disease; rheumatoid arthritis; idiopathic pulmonary fibrosis; angiogenesis-dependent chronic inflammatory condition; lung cancer; melanoma; inflammatory disease; diabetes; arteriosclerosis; cataract; reperfusion injury; cancer; meningitis; rheumatic disease; skin disease; idiopathic pulmonary fibrosis; inflammatory bowel disease; psoriasis; ulcerative colitis; eczema; atopic dermatitis; contact dermatitis; inflammatory lung disease; ARD; adult respiratory disease; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The sequence is a Human interleukin-20, IL-20 protein of the invention. The invention relates to treating a mammal afflicted with a disease in which an interleukin-20 (IL-20) polypeptide plays a role, involves administering antagonist of IL-20 polypeptide to the individual. The method is useful for treating psoriasis, eczema, atopic dermatitis, contact dermatitis, adult respiratory disease, asthma, bronchitts and pneumonia and is also useful for treating multiple organ failure, inflammatory lung injury, septic shock, bacterial pneumonia, inflammato bowel disease, rheumatoid arthritis, ulcerative colitis and Crohn's
                                                                                    (FOST/)
                (SPRE/)
(BLUM/)
(EAGA/)
                                                                                                                                                         23-DEC-1999;
                                                                                                                                                                                    22-DEC-2000; 2000US-00746359.
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                                                                                                                                                                                                                                                                                                       bronchitis;
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                                                                                                                                             22-JUN-2000;
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176; Conserv
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XU W.

MADDEN K L.

KELLY J D.

SPRECHER C A.

BLUMBERG H.

EAGAN M A.

JASPERS S R.
                                                                                                 THOMPSON
FOSTER D
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                                                                                                                                                                                                                                                                                                       pneumonia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                disease related human interleukin (IL)20 polypeptide
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                                                                                                                                             2000US-0213341P
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Pred. No. 5e-88;
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melanoma cell lines. Therefore an effective method to treat inflammatory diseases would be to administer an agent that would inhibit II-8. It has been shown that II-20 up-regulates II-8. Therefore antagonists to II-20 can be used to treat these diseases. The method is used for treating diseases in which the II-20 polypeptide plays a role e.g. inflammatory diseases including diabetes, arteriosclerosis, cataracts, reperfusion injury, cancer, meningitis, rheumatic disease, idiopathic pulmonary fibrosis, inflammatory bowel disease (ulcerative colitis), skin disease (psoriasis, eczema, atopic dermatitis and contact dermatitis) or an inflammatory lung disease (adult respiratory disease (ARD), asthma, bronchitis and pneumonia). This sequence represents a human interieukin-
                                                                                                                                                                                                                                                                                                         receptors on neutrophils, monocytes, basophils, and eosinophils. IL-8 a potent chemoattractant for neutrophils; and the early stages of periodontal disease are characterised by the influx of neutrophils. IL-is a potent inducer of angiogenesis in several angiogenesis-dependent chronic inflammatory conditions, including rheumatoid arthritis, psoriasis, and idiopathic pulmonary fibrosis. Additionally, IL-8 is an important source of angiogenic activity in human lung cancer. Also, IL-expression correlates with experimental metastatic activity of some
                                                                                                                                                                                                                                                                                                                                                                                                                                                                comprising administering antagonist of the II-20 polypeptide plays a role individual. An important cytokine in the inflammatory process is interleukin-8 (II-8), a chemokine that acts as an agonist for neutrophils via chemotaxis and the release of granule enzymes. II-8 binds to receptors on neutrophils, monocytes hasoning.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Thompson
Blumberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 12-13; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treating inflammatory skin and lung diseases using antibodies against interleukins (IL)-20 (which indirectly modulates activation of IL-8),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-507215/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CHAN/) CHANDRASEKHER Y (NOVA/) NOVAK J E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     he invention describes a method (I) disease in which an interleukin-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for treating e.g. psoriasis, asthma and bronchitis.
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                                       polypeptide used
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Eagan MA,
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Jaspers
                                       in developing
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SR, Ch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for treating a mammal afflicted with (IL-20) polypeptide plays a role
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  en KL, Kelly JD,
Chandrasekher YA,
                                            the method
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Novak JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            against
                                                                                                                                                                                                                                                                                                                                     IL-8
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RESULT 5
AAE29052
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Best Local Sim
Matches 176;
 Human
                        27-JAN-2003
                                               AAE29052;
                                                                   AAE29052 standard;
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 IL-20
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protein
                        (first
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밁 S S 밁 Ś

Sequence 176 AA;

Similarity

100.0%;

Score 910; DB 9

5

Length 176;

Conservative

TXXXXXXXXXX

Human; heterodimeric cytokine receptor; interleukin-22R; IL-22R; asthma; inflammatory disease; psoriasis; adult respiratory disease; bronchitis; septic shock; multiple organ failure; inflammatory lung injury; eczema;

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RESULT 6
ABR62464
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Best Local
                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to novel heterodimeric cytokine receptor which comprises an interleukin-22R (II-22R, ZcytoRII)) subunit. Receptor sequences are useful for down-regulating II-20 and treating inflammatory diseases such as psoriasis, adult respiratory disease, multiple organ failure, septic shock, inflammatory lung injury such as bronchitis or asthma, bacterial pneumonia, eczema, atopic and contact dermatitis, ulcerative colitis and Crohn's disease. The present sequence is human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Soluble heterodimeric cytokine receptor useful for interleukin-20 and treating inflammatory diseases, asthma, comprises an interleukin-22R subunit and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bacterial pneumonia; dermatitis; ulcerative colitis; Crohn's disease; antiinflammatory; dermatological; antibacterial; immunosuppressive;
                                                                Human;
                                                                                     Human
                                                                                                          06-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-723314/78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chandrasekher YA,
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21-JUN-2001;
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                                           Homo sapiens
                                                                                                                                                  ABR62464
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            Peptide
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176;
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                                                                interleukin-20;
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                                                                                                                                                standard; protein; 176
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                                                                                                                                                                                                                                                                                                                               Conservative
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2001US-0299865P
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1. .24
/label= Signal_peptide
                      Location/Qualifiers
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                                                                                                         entry)
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                                                                IL-20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English.
                                                                                                                                                                                                                                                                                                                               0
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Pred. No. 5e-88;
; Mismatches 0;
                                                                cervical
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                                                                cancer;
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                                                                cytostatic;
                                                                                                                                                                                                                                                                                                                                                  Length 176;
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                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     down-regulating
such as psoriasis
interleukin-20RB
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                                                                 virucide
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                                                                                                                                                                                                                          RESULT 7
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Query Match
                  Key
Peptide
                                                                                                      autoimmune disease; insulin-dependent unaweves muration; radiation therapy; rheumatoid arthritis; multiple sclerosis; infection; radiation therapy; asthma; bronchitis; wound healing; periodontal disease; antiinflammatory; nsoriasis; eczema; dry skin; oral cavity; muscle atrophy; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                virus. IL-20 can be or chemotherapeutic lesions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is the protein sequence of human interleukin-20 (IL-20), including the signal sequence. The invention relates to the use of IL-20 for treating cervical cancer or cells infected with human papilloma virus. IL-20 can be administered alone or in conjunction with radiation or chemotherapeutic agents or surgical excision of the involved cells or legicar.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Inhibiting the growth and/or proliferation of cervical cancer cells, useful for treating cervical cancer or human papilloma virus infection, comprises bringing Interleukin-20 (IL-20) into contact with the cervica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-DEC-2002; 2002WO-US040309
                                                                                                   cancer;
                                                                                                                                                                                      Human;
                                                                                                                                                                                                                   Human
                                                                                                                                                                                                                                                                                    ABW00889;
                                                                                                                                                                                                                                                                                                                    ABW00889 standard;
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                                                                  Homo sapiens
                                                                                                                                                                                                                                                    15-JAN-2004
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)B; ACF05161.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  176;
                                                                                                                                                                                                                 cytokine-like polypeptide-10 (Zcyto10)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61
                                                                                                                                                                                   cytokine-like
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                                                                                                    vulnerary.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                KDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMEETE 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                DIRILRRTESLQDTKPANRCCLLRHLLRLYLDRVFKNYQTPDHYTLRKISSLANSFLTIK
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Location/Qualifiers
1. .24
/label= Signal_peptide
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/label= IL-20
                                                                                                                                                                                                                                                                                                                   protein;
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                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                  polypeptide-10; Zcyto10; therapy; thrombocytopaenia; insulin-dependent diabetes mellitus; immune response;
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                                                                                                                                                                                                                                                                                                                    176
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Pred. No. 5e-88;
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Protein

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                                                                                                                                                                                                                                                                                                                           The invention relates to an isolated human cytokine-like polypeptide-10 CC (Zcytol0) polypeptide. Zcytol0 is useful for prevention or treatment of CC conditions characterised by improper cell proliferation, cell CC differentiation or cytokine production. Zcytol0 is useful for treating CC rheumatoid arthritis, multiple sclerosis, etc., by inhibiting calcular companies. The invention is useful for inhibiting cancer cell component or proliferation, for stimulating immune system to combat CC microbial or viral infections, for increasing platelet production, and thus useful in cancer patients who experience thrombocytopaenia due to cancer therapy or radiation therapy. Zcytol0 is useful in tracheacy specific or tracheabronchial specific applications, such as in the condition or radiation therapy. Zcytol0 is useful in tracheacy cunderlying the same, in regulating mucous production or mucociliary conditions or in treatment of asthma, bronchitis or other CC diseases of the tracheabronchial tract. Zcytol0 may also enhance wound cells consistent of the same, in regulating mucous production may be capacially useful in the treatment of affected tissues which may be capacially useful in the treatment of periodontal disease. Zcytol0 can be cused to treat skin conditions such as psoriasis, eczema and dry skin in CC capacially and for modulating muscle tone in the tracheabronchial tract, and for treating muscle atrop tone in the tracheabronchial tract, condition useful for promoting wound healing. The present sequence cis human Zcytol0 protein
                                                                                                                                                                                                            Query Match
Best Local S
Matches 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel human cytokine-like polypeptide-10, Zcyto10, useful for treating insulin-dependent diabetes mellitus, rheumatoid arthritis, multiple sclerosis, asthma, psoriasis and inhibiting cancer cell growth.
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25-NOV-1998;
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 121
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                    KDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMEETE
                                                                                                                                                         MKASSIAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNI
KDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMEETE
                                                                    DIRILRRIESLODTKPANRCCLLRHLLRLYLDRVFKNYQTPDHYTLRKISSLANSFLTIK
                                                                                          DIRILRRTESLQDTKPANRCCLLRHLLRLYLDRVFKNYQTPDHYTLRKISSLANSFLTIK
                                                                                                                                      MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNI
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98US-00199586.
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/note= "M
33. .126
80. .132
81. .134
                                                                                                                                                                                                                                                                                                                    protein
                                                                                                                                                                                                                           100.0%; Score 910; DB 7 100.0%; Pred. No. 5e-88;
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                                                                                                                                                                                                                                              DB 7;
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                                                                                                                                                                                                                                            Length 176;
                                                                                                                                                                                                            Indels
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176
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Best Local Sin Matches 176;

Similarity

Conservative

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Gaps

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DIRILRRTESLQDTKPANRCCLLRHLLRLYLDRVFKNYQTPDHYTLRKISSLANSFLTIK 120

MKASSIAFSILSAAFYILWTPSTGLKTINIGSSCVIATNIQEIRNGFSDIRGSVQAKDGNI MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNI

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ADG46669
                 Query Match
Best Local
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25-NOV-1998;
17-MAY-1999;
                                                                                                      regulating mucous production; mucocilary clearance of debris; in treatment of asthma, bronchitis and other diseases of the tracheobronchial tract. They are also used to promote regeneration of affected tissues which may be especially useful in the treatment of periodontal disease. Zcytolo polypeptides can be used to treat skin conditions such as psoriasis, eczema and dry skin in general. They are also used in protein therapy. The present sequence is human Zcytolo
                                                                                                                                                                                                                 The present invention relates to novel mammalian cytokine-like polypeptide-10 (Zcyto10) proteins and polynucleotides encoding such proteins. Sequences of the invention are useful for promoting the he of wounds and for stimulating the proliferation of platelets. In particular Zcyto10 polypeptides may be used in trachea-specific or tracheobronchial-specific applications such as maintenance or wound repair of the tracheobronchial epithelium or cells underlying it; in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cytokine-like polypeptide-10; Zcyto10; wound healing; platelet proliferation; wound repair mucous production; asthma; bronchitis; tracheobronchial tract disease; periodontal disease; skin condition; psoriasis; eczema; dry skin; protein therapy; hu vulnerary; respiratory; antibronchitic; dental; dermatological.
                                                              Sequence 176 AA;
                                                                                                                                                                                                                                                                                                                                                                         Novel isolated mammalian cytokine-like polypeptide-10 (Zcyto10), useful for treating asthma, bronchitis and other tracheobronchial damage.
                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 2; 31pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB; ADG46668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conklin DC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-APR-2003; 2003US-00413661
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CONK/) CONKLIN D C. (HALD/) HALDEMAN B A.
                                                                                                                                                                                                                                                                                                                                                                                                                                        2003-863865/80.
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98US-00199586.
99US-00313458.
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                 100.0%;
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Score 910; DB 7
Pred. No. 5e-88;
; Mismatches
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                                DB 7;
                                Length 176;
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                                                                                                                                                                                                                                                                               the healing
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RESULT 9
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22-JUN-2000;
22-DEC-2000;
         The invention relates to a novel method of reducing or treating inflammation in a mammal which comprises administering an antibody, antibody fragment or single chain antibody which specifically binds to a receptor of interleukin (II)-20 comprising an II-20RA subunit and an II-20RB subunit. The method of the invention has antiinflammatory, dermatological, antipsoriatic, antiarthritic, respiratory, antiasthmatic, antiulcer, antibacterial, immunosuppressive and gastrointestinal applications and may be useful for reducing or treating an inflammation, including an inflammatory skin disease such as psoriasis, eczema, atopic including an inflammatory skin disease such as psoriasis.
                                                                                                                                                                                                                                                      Thompson
Blumberg
                                                                                                                                                               Reducing or treating inflammation, e.g. inflammatory lung disease, comprises administering an antibody, antibody fragment or single chain antibody that specifically binds to an interleukin (IL)-20RA subunit of an IL-20 receptor.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US2004005320-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ulcerative colitis; Crohn's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           atopic
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antiarthritic; respiratory; antiasthmatic; antiulcer; antibacterial;
immunosuppressive; gastrointestinal; skin; psoriasis; eczema;
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(FOST/)
(XUWW/)
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                                                                                                                                        Disclosure;
                                                                                                                                                                                                                                WPI; 2004-081696/08.
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                                                                                                                                                                                                                                                                                                                                                           SPRE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              c dermatitis; contact dermatitis; lung; respiratory distress syndrome; asthma; bronchitis; itis; septic shock; multiple organ failure; bowel;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61
                                                                                                                                                                                                                                                                                                       EAGAN M A.
JASPERS S R.
CHANDRASEKHER
                                                                                                                                                                                                                                                                                                                                            KELLY J D.
SPRECHER C A.
BLUMBERG H.
                                                                                                                                                                                                                                                                                             NOVAK J
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                                                                                                                                                                                                                                                      Foster DC,
Eagan MA,
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2000US-0213341P.
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                                                                                                                                                                                                                                                      Madden KL, Kelly JD,
SR, Chandrasekher YA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             interleukin; IL-20; IL-20RA subunit;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ij
lung disease
                                                                                                                                                                                                                                                       Sprecher CA;
Novak JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pneumonia
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inflammatory

Query Match Best Local S Matches 176

Local Similarity nes 176; Conserv

100.0%; ilarity 100.0%; Conservative 0

0,

Score 910; DB 8 Pred. No. 5e-88; ; Mismatches

8

Length 176;

Indels

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Gaps

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Sequence

176

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RESULT 10
ADM95041
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                    This invention relates to a novel agent for prevention and/or treatment of colon cancer, comprising of the interleukin 20 (II-20) receptor alpha chain, its partial peptide or salt. The invention may be useful for the development of compounds with a cytostatic activity. The invention is useful for preventing and/or treating colon cancer. A diagnostic agent is useful for diagnosis of colon cancer. By using II-20, II-20 receptor, its salt or its peptide, compounds that have altered binding property towards them can be efficiently screened. The present sequence is that of the human Interleukin-20 protein which is related to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               colon
IL-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adult respiratory distress syndrome, asthma, bronchitis and pneumonia, well as arthritis, septic shock, multiple organ failure, inflammatory bowel disease, ulcerative colitis or Crohn's disease. The current sequence is that of the human IL-20 protein of the invention.
                                                                                                                                                                                                                                                WPI; 2004-320878/30.
N-PSDB; ADM95042.
                                                                                                                                                                                                         Agent useful
interleukin :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADM95041 standard;
                                                                                                                                                                  Disclosure;
                                                                                                                                                                                                                                                                                                                     12-AUG-2002; 2002JP-00234880
                                                                                                                                                                                                                                                                                                                                                                             11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                         JP2004075569-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-JUN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor alpha chain; cytostatic; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Interleukin-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer; interleukin 20 receptor alpha chain;
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                                                                                                                                                                                                                                                                                            TAKEDA CHEM
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                                                                                                                                                                                                        l for prevention and/or treatment
20 (IL-20), IL-20 receptor alpha
                                                                                                                                                                 SEQ ID NO 9;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first
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                                                                                                                                                               50pp; Japanese.
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Pred. No. 5e-
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8
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                                                                                                                                                                                                         of colon cancer, comprises chain, IL-20 receptor beta
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cell differentiation; cytokine production; autoimmune disease; insulin dependent diabetes mellitus; multiple sclerosis; rheumatoid arthritis; cancer; wound healing; tissue regeneration; platelet count; asthma; skin disease; psoriasis; eczema; dry skin; antidiabetic; neuroprotective; antirheumatic; antiarthritic; cytostatic; antidiabetic; recognication;
The present invention provides mammalian cytokine-like (Zcyto10) proteins and their encoding polynucleotides.
                                               Claim 1; SEQ ID NO 2; 31pp; English.
                                                                                             New Zcyto10 polypeptides and polynucleotides, useful for treating autoimmune diseases (e.g. insulin dependent diabetes mellitus, multiple sclerosis or rheumatoid arthritis), or skin conditions (e.g. psoriasis or
                                                                                                                                                                 N-PSDB;
                                                                                                                                                                                                                                                                                            26-NOV-1997;
25-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disulfide-bond
                                                                                                                                                                                                               Conklin
                                                                                                                                                                                                                                                                                                                                             27-FEB-2004; 2004US-00789129
                                                                                                                                                                                                                                                                                                                                                                              22-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mammalian cytokine-like polypeptide-10; Zcyto10; cell proliferation; cell differentiation; cytokine production; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human Zcyto10 longer form protein.
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                                                                                                                                                                                                                                                                                15-APR-2003;
                                                                                                                                                                                                                                                 ( OMYZ)
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                                                                                                                                                                               2004-552564/53
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                                                                                                                                                                                                               DC,
                                                                                                                                                                                                                                                ZYMOGENETICS INC.
                                                                                                                                                                ADQ88066
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                                                                                                                                                                                                               Haldeman
                                                                                                                                                                                                                                                                                2003US-00413661.
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                                                                                                                                                                                                                                                                                               97US-0066597P.
98US-00199586.
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25. .176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note = Intramolecular disulfide bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                              /note = Intramolecular disulfide bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               . 134
                                                                                                                                                                                                               BA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Human mature Zcytol0 protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Intramolecular disulfide bond
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The invention
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WPI; 2004-580197/56

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Best Local 9
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Disulfide-bond
Disulfide-bond
                                                                                                                                                                        26-NOV-1997;
25-NOV-1998;
15-APR-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               useful for treating conditions characterised by improper cell proliferation, cell differentiation or cytokine production, specifically for treating autoimmune diseases such as insulin dependent diabetes mellitus, multiple sclerosis and rheumatoid arthritis and for inhibiting cancer cell growth. The invention is also used to stimulate the immune system, to enhance wound healing and promote regeneration of affected tissues, to increase platelet count and to treat asthma and skin conditions such as psoriasis, eczema and dry skin. The invention acts as an antidiabetic, neumostimulant, vulnerary, antiasthmatic, antipsoriatic and dermatological agent. The invention is also used in gene therapy. The present sequence is human Zcytol0 longer form protein.
                                                                        Conklin
                                                                                                                          OMYZ)
                                                                                                                                                                                                                                                                         27-FEB-2004;
                                                                                                                                                                                                                                                                                                                             05-AUG-2004.
                                                                                                                                                                                                                                                                                                                                                                           US2004152878-A1
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psoriasis; eczema; dry skin; w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human
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                                                                                                                            ZYMOGENETICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNI
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                                                                                                                                                                      ; 97US-0066597P.
; 98US-00199586.
; 2003US-00413661.
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                                                                                                                                                                                                                                                                           2004US-00789968
                                                                           Haldeman
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                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "M
33. .126
80. .132
81. .134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1. .24
/label= Signal_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polypeptide-10 (Zcyto10) long form protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
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                                                                                                                            INC
                                                                           BA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Mature Zcyto10 long
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Zcyto10; therapy; asthma; infection; wound healing; platelet proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 910; DB 8
Pred. No. 5e-88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ₽
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein"
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RESULT 13
ADS18363
ID 18363
ID 18363
ID 2018
XX ADS18
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zcyto10; cytokine-like polypeptide; cell proliferation; cell differentiation; autoimmune disease; diabetes; multiple scl rheumatoid arthritis; asthima; psoriasis; cancer; wound; gene the immunosuppressive; antidiabetic; neuroprotective; antirheumatic; antiarthritic; antiasthmatic; antipsoriatic; cytostatic; vulnera
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated antibody that binds to a Zcytol0 polypeptide, useful treating asthma, microbial or viral infections, and for promoting healing of wounds, psoriasis, eczema or dry skin.
    Region
                                                                                                                                                                                                                                                                       Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  proliferation
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                                                                                                                                                                                                                                                                                                                   Region
                                                                                                                                                                                                                                                                                                                                        Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                          Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention relates to novel cytokine-like polypeptide-10 (Zcyto10) ypeptides and polynucleotides. Zcyto10 sequences are useful for ating asthma, microbial or viral infections, psoriasis, eczema or n, for promoting the healing of wounds and for stimulating the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMEETE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DIRILRRTESLODTKPANRCCLLRHLLRLYLDRVFKNYQTPDHYTLRKISSLANSFLTIK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNI
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33. .126
35
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                                                                                                                                                                                                                                                                                                                                                                                 /label= Signal_peptide
25. .176
      /note=
126
/note=
158
/note=
172
                                                                                                                                                                                /note=
105
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                                                                                                                                                                                                       "Helix B"
                                                                                                                                                                                                                                                                                              "Helix
                                                                                                                                                                                                                                                                                                                                                              "Mature Zcytol0 protein"
                             "Helix D'
                                                                         "Helix
                                                                                                                    "Helix C"
                                                                                                                                                              "Helix
                                                                                                                                                                                                                                                   "Helix
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 910; DB 8
Pred. No. 5e-88;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human Zcytol0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              176
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RESULT 14
ADV66904
ID ADV66
XX ADV66
XX ADV66
XX ADV66
XX ADV66
XX DOT 24-FE
DT 24-FE
DT Humar
XX Derma
KW Derma
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                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to a mammalian cytokine-like polypeptide, Zcytol0 and its encoding polynucleotide. The invention is useful for modulating cell proliferation, cell differentiation or cytokine production in the prevention or treatment of conditions such as autoimmune diseases (e.g. diabetes, multiple sclerosis, rheumatoid arthritis, asthma or psoriasis) and cancer or wounds. The invention is also useful in gene therapy. The present sequence is the human Zcytol0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antibody that selectively binds to a ZcytolO polypeptide useful for modulating cell proliferation, cell differentiation or cytokine production in the prevention or treatment of e.g. autoimmune diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-NOV-1997; 97US-0066597P
25-NOV-1998; 98US-00199586
15-APR-2003; 2003US-00413661
                                                                                                                            24-FEB-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer or wounds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-DEC-2003; 2003US-00748484
                                                           interleukin-20;
                                                                        Dermatological;
                                                                                                   Human
                                                                                                                                                     ADV66904;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein.
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DB; ADS18362.
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                                                                                                                                                                                                                                                                                                                                                                                               176;
                                                                                                 IL-20 wild-type, SEQ
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HALDEMAN B A.
GROSSMANN A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DC,
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                                                                                                                                                                             standard;
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                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
llarity 100.0%;
Conservative
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                                                              Antiinflammatory;
IL-20; cytokine;
                                                                                                                                                                           protein;
                                                                                                                            entry)
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                                                            cytokine;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English
                                                                                                                                                                              176 AA
                                                                                                                                                                                                                                                                                                                                                                                              Score 910; DB 8
Pred. No. 5e-88;
; Mismatches
                                                                                                   20
                                                              ; Interleukin-20 antagonist; dermatological disease; inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                       8
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                                                                                                                                                                                                                                                                                                                                                                                                                       176;
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US2004248097-A1

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ADW64518
ID ADW64
XX ADW64
XX ADW64
XX ADW64
XX ADW64
XX Thera
XX Thera
KW Antii
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Matches 176
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                                  gastrointestinal
                                                Therapy; cns-gen.; respiratory-gen.; antiasthmatic; antipsoriatic; antiinflammatory; psoriasis; dermatological disease; immune disord pulmonary disease; asthma; inflammation; respiratory disease; bron antiinflammatory; inflammation; cystic fibrosis;
                                                                                                                                       Human
                                                                                                                                                                                                                                            ADW64518
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                                                                                                                                                                         24-MAR-2005
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20 variants
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                                disease; chemotherapy; interleukin.
                                                                                                                                                                                                                                          protein; 176
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Matches 176;
                                                                                                                                                                                            The present invention relates to an soluble receptor having an interleukin 20 receptor alpha (IL-20RA) subunit and an IL-20RB subunit. The invention is useful for treating inflammatory diseases such as psoriasis and inflammatory lung diseases such as asthma, bronchitis and cystic fibrosis. The present sequence is the human IL-20 protein.
                                                                                                                                                                                                                                                                                               New isolated soluble receptor comprised of an interleukin 20 receptor alpha (IL-20RA) subunit and a IL-20RB subunit, useful for treating inflammatory diseases e.g., psoriasis and inflammatory lung diseases
                                                                                                                                                                            Sequence 176 AA;
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                                                                                                                                                                                                                                                                 Disclosure;
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22-JUN-2000; 2000US-0213416P
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1 MKASSLAFSLLSAAFYLLWT.....AVVKALGELDILLQWMEETE 176
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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A;Gene: GDB:IL10; IL-10 A;Gene: GDB:IL10; IL-10 A;Gross-references: GDB:128636; OMIM:124092 A;Map position: 1931-1932 A;Introns: 55/3; 75/3; 126/3; 148/3 A;Introns: 55/3; 75/3; 126/3; 148/3 C;Superfamily: interleukin-10	A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-6 <pla> A;Cross-references: UNIPARC:UPI000011E9B9; EMBL:X73536; NID:g452395; PIDN:CAA51942.1; EC.Comprise:</pla>	A; Molecule type: DNA A; Residues: 1-6, PVAMSS' <kub> A; Cross-references: UNIPARC: UPI0000178477; EMBL: X78437; NID: 91167482 A; Cross-references: UNIPARC: UPI0000178477; EMBL: X78437; NID: 91167482 R; Platzer; C.; Volk, H.D.; Platzer, M. DNA Seg. 4, 399-401, 1994 A; Title: 5' noncoding sequence of human IL-10 gene obtained by oligo-cassette PCR walk- A; Reference number: I37890; MUID: 95143580; PMID: 7841462 A; Accession: I37890; MUID: 95143580; PMID: 7841462 A; Accession: I37890; MUID: 95143580; PMID: 7841462</kub>	A;Status: translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-178 <52Nn> A;Residues: 1-178 <52Nn> A;Residues: 1-178 <52Nn> A;Residues: 1-178 <52Nn A;Residues: 1-178 <52Nn A;Residues: 1-178 <52Nn A;Cross-references: UNIPARC:UPIO000034E50; EMBL:U16720; NID:g1041812; PIDN:AAA80104.1; R;Kube, D.; Platzer, C.; von Knethen, A.; Straub, H.; Hafner, M.; Tesch, H. submitted to the EMBL Data Library, March 1994 A;Description: Isolation of the human interleukin-10-promoter. Characterization of the A;Reference number: S49110 A;Accession: S49110 A;Accession: S49110 A;Status: preliminary	A;Accession: A38580 A;Molecule type: mRNA A;Residues: 1-178 «VIE» A;Residues: 1-178 «VIE» A;Cross-references: UNIFROT:P22301; UNIFARC:UPI0000034E50; GB:M57627; NID:g186270; PID: A;Cross-references: UNIFROT:P22301; UNIFARC:UPI0000034E50; GB:M57627; NID:g186270; PID: A;Windsor, W.T.; Syco, R.; Tearbopoulos, A.; Zhang, R.; Durkin, J.; Baldwin, S.; Paliw. Biochemistry 32, 8807-8815, 1993 A;Title: Disulfide bond assignments and secondary structure analysis of human and muri. A;Reference number: A48693; MUID:93372085; PMID:8364028 A;Contents: annotation; disulfide bonds in recombinant protein R;Sanjanwala, B. R;Sanjanwala, B. R;Sanjanwala, B. A;Reference number: G07695 A;Accession: G01539	RESULT 1 A38580 interleukin-10 precursor - human interleukin-10 precursor - human N;Alternate names: cytokine synthesis inhibitory factor (CSIF); IL-10 C;Species: Homo sapiens (man) C;Date: 14-Feb-1992 #sequence revision 14-Feb-1992 #text_change 09-Jul-2004 C;Accession: A38580; G01539; S49110; I37890 R;Vieira, P.; de Waal-Malefyt, R.; Dang, M.N.; Johnson, K.E.; Kastelein, R.; Fiorentin Proc. Natl. Acad. Sci. U.S.A. 88, 1172-1176, 1991 A;Title: Isolation and expression of human cytokine synthesis inhibitory factor cDNA c A;Reference number: A38580; MUID:91142134; PMID:1847510	ALIGNMENTS

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C;Keywords: cytokine; glycoprotein; lymphokine; T-cell F;1-18/Domain: signal sequence #status predicted <SIG> F;19-178/Product: interleukin 10 #status predicted <MAT> F;30-126,80-132/Disulfide bonds: #status experimental F;134/Binding site: carbohydrate (Asn) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Title: DNA sequence and expression of the B95-8 Epstein-Barr virus genome. A;Reference number: A03794; MUID:84270667; PMID:6087149
A;Contents: annotation; protein coding region
C;Superfamily: interlankin-10
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                        R:Blancho, G.; Gianello, P.; Germana, S.; Baetscher, M.; Sachs, D.H.; LeGuern, Proc. Natl. Acad. Sci. U.S.A. 92, 2800-2804, 1995
A;Title: Molecular identification of porcine interleukin 10: regulation of exp A;Reference number: I46591; MUID:95224028; PMID:7708727
A;Accession: I46591
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Accession: A03741
A;Molecule type: DNA
A;Residues: 1-170 <BAN>
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A;Title: Sequence analysis of the 17,166 bp EcoRI fragment C o A;Reference number: A93065; MUID:85035713; PMID:6092825
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;Species: Sus scrofa domestica (domestic pig)
;Date: 21-Feb-1997 #sequence_revision 21-Feb-1997 #text_change
;Accession: I46591
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STGLKTLNLGSCV----IATNLQEIRNGFSDIRGSVQAKDGNIDIRILRRTESLQDTKP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QTPDHYTLRKISSLANSFLTIKKDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAV 158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOBIRNGFSDIRGSVOAKDGNIDIRILRRTESLQDTKPANRCCLLRHLLRLYLDRVFKNY
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                                                                                                                                                                                                                                                                                                                               YKAMSEFDIFINYIE
                                                                                                                                                                                                                                                                                                                                                                        VKALGELDILLOWME 173
                                                                                                                                                                                                                                                                                                                                                                                                                          ENQDPEAKDHVNSLGENLKTLRLRRCHRFLPC---ENKSKAVEQIKNAFNKLQ-EKGI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LRDLRDAFSRVKTFFQTKD-EVD-NLLLKESLLEDFKGYLGCQALSEMIQFYLEEVMPQA
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27.4%;
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evision 25-Feb-1985
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Pred. No. 8.1e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 155; DB 1;
Pred. No. 1.1e-07;
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                                                                          expression
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RESULT 5 A34853 interleukin-10]

precursor -

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interleukin 10 homolog - equine herpesvirus 2
N;Alternate names: hypothetical protein E7
C;Species: equine herpesvirus 2
C;Date: 17-Feb-1994 #Bequence revision 17-Feb-1994 #text_change 09-Jul-2004
C;Accession: A48558; S55664
R;Rode, H.J.; Jansen, W.; Rosen-Wolff, A.; Bugert, J.J.; Thein, P.; Becker, Virus Genes 7, 111-116, 1993
A;Title: The genome of equine herpesvirus type 2 harbors an interleukin 10 (1)
A;Reference number: A48558; MUID:93227565; PMID:8385838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: mRNA
A;Residues: 1-175 <BLA>
A;Cross-references: UNIPROT:Q29055;
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPROT:Q89451; UNIPARC:UPI000012D058; GB:S59624; NID:g299468; A;Note: sequence extracted from NCBI backbone (NCBIN:129334, NCBIP:129335)
A;Rote: fequence extracted from NCBI backbone (NCBIN:129334, NCBIP:129335)
A;Telford, E.A.R.; Watson, M.J.
J. Mol. Biol. 249, 520-528, 1995
A;Title: The DNA sequence of equine herpesvirus 2.
A;Reference number: S55594; MUID:95302501; PMID:7783207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Gene: I1-10
C; Superfamily:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Reference number: A; Accession: A48558
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A; Residues: 1-179 < TEL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Accession: S55664
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A; Residues: 1-179 < ROD>
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                                                                                                                                                                                                                                                                                                                                                                                Cross-references: UNIPARC:UPI000012D058; GB:U20824; NID:g695172; PIDN:AAC13857.1; Note: the nucleotide sequence was submitted to the EMBL Data Library, February 19;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    116 FLTIKKDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 MPSSALLYCLI-----FLAGVAASIKSEN--SCIHFPTSLPHMLRELRAAFGPVKSFFQT 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCV----IATNLQEIRNGFSDIRGSVQA
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                                                                                                                                                                                                                                                                                            Similarity
                                         AAVVKALGELDILLQWME
                                                                                      QAENHSTDQEKDKVNSLGEKLKTLRVRLRRCHRFLPC---ENKSKAVEQVKSAFSKLQ-E
                                                                                                                                                                              LHELRAAFSRVKTFFOMKD-QLD-NMLLDGSLLEDFKGYLGCQALSEMIQFYLEEVMP--
                                                                                                                                                                                                                      LOEIRNGFSDIRGSVQAKDGNIDIRILRRTESLQDTKPANRCCLLRHLLRLYLDRVFKNY
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KGVYKAMSEFDIFINYIE 170
                                                                                                                                  QTPDHYTLR---KISSLANSFLTIKKDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQ 155
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27.0%;
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                                                                                                                                                                                                                                                                  24;
                                            173
                                                                                                                                                                                                                                                                     Score 147.5; DB 2
Pred. No. 6.1e-07;
4; Mismatches 59
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                               DB 2;
                                                                                                                                                                                                                                                                       59;
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R; Feng, L.; Tang, W.W.; Chang, J.C.C.; Wilson, Biochem. Biophys. Res. Commun. 192, 452-458, 19 A; Title: Molecular cloning of rat cytokine synt A; Reference number: JN0475 A; Accession: JN0475 A; Accession: JN0475 A; Status: nucleic acid sequence not shown A; Molecule type: mRNA A; Residues: 1-178 < FEND A; FERSIDES: TITLES: TOTALE: TENDOT: DOLGS: 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N;Alternate names: cytokine synthesis inhibitory factor C;Species: Rattus norvegicus (Norway rat) C;Date: 30-Sep-1993 #sequence revision 30-Sep-1993 #text_change C;Accession: JN0475; JC1357; $\overline{8}36021
                                                                                                                                A;Molecule type: mRNA
A;Residues: 1,'P',3-64,'L',66-178 <GOO>
A;Cross-references: UNIPARC:UPI000012D477; GB:L02926;
                                                                                                                                                                                                                                              A; Reference number: A; Accession: JC1357
                                                                                                                                                                                                                                                                       R;Goodman, R.E.; Oblak, J.; Bell, R.G.
Biochem. Biophys. Res. Commun. 189, 1-7, 1992
A;Title: Synthesis and characterization of rat
A;Reference number: JC1357; MUID:93080542; PM
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C;Superfamily: interleukin-10
C;Keywords: cytokine; lymphokine; T-cell
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A;Title: Structure of the mouse II-10 gene and chromosomal A;Reference number: I56136; MUID:9226850B; PMID:1350294
A;Accession: I56136
A;Reference 1
A;Accession:
                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPROT: P29456; UNIPARC: UPI0000167959
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A; Residues: 1-178
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                             the EMBL Data Library, July 1991
number: S36021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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9; Mismatches
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                                                                                                                                   NID:g204903; PIDN:AAA41425.1;
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A;Introns: 1/3; 38/3; 81/1; 143/1; 211/1; 238/2; 278/3; 300/3; 336/2;
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A;Residues: 1-567 <SON>
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Biochem. Biophys. Res. Commun. 235, 10-14,
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                                                                                                                                                                                                                                                                                                                                       enzyme
                                                                                                                                                                                                                                                                                              383/2;
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RESULT

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A45977
Rab geranylgeranyl transferase component B alpha subunit - rat C.Species: Rattus norvegicus (Norway rat)
C.Date: 16-Feb-1994 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C.Accession: A45977
R;Armstrong, S. A.; Seabra, M.C.; Sudhof, T.C.; Goldstein, J.L.; Brown, M.S.
J. Biol. Chem. 268, 12221-12229, 1993
J.Title: cDNA cloning and expression of the alpha and beta subunits of rat Rab. A;Reference number: A45977; MUID:93280201; PMID:8505342
A;Accession: A45977
A;Bolecule type: mRNA; protein
                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:P46682; UNIPARC:UPI00000530A8; A;Experimental source: strain S288C R;Robinson, L.C.; Engle, H.M.; Panek, H.R. submitted to the EMBL Data Library, September 1995 A;Description: Suppressors of loss of yeast casein kinase 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       probable beta-adaptin YKS5 - yeast (Saccharomyces cerevisiae)

N;Alternate names: protein G9331; protein YGR261C

C;Species: Saccharomyces cerevisiae

C;Date: 17-May-1996 #sequence_revision 17-May-1996 #text_change 09-Jul-2004

C;Accession: S64594; S63450

R;Panzeri, L.; Agostoni Carbone, M.L.; Melchioretto, P.; Plevani, P.; Martegani, submitted to the Protein Sequence Database, May 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT
S64594
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                                    á
                                                                                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-26, 'PLSWVNPP', 36-723, 'T', 725-795, 801-809 .
A; Cross-references: UNIPARC: UPI0000168A41; EMBL: U35411;
                                                                                                                                                                                                                                                                                                                                 A; Accession: S63450
                                                                                                                                                                                                                                                                                                                                                     A; Description: Suppressors A; Reference number: S63439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: DNA
A; Residues: 1-809 < PAN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Reference number: S64591
A; Accession: S64594
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Note: sequence extracted from NCBI backbone (NCBIN:133369, NCBIP:133370
                                                                                                                                                                             ;Map
                                                                                                                                                                                                 Cross-references: SGD:S0003493; MIPS:YGR261c
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Best Local :
                                                                                      Matches
                                                                                                                                                                             position:
                                                                                                             Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EKSTVLQSELESCKELQELEPENKWCLLTIILLMRALDPLLYEKETLQYFSTLKAVDPMR
EAAAVATSKLGESSYTYY--SQNINPQQLVTLLNSRNSREVRDAMKRIISIMASDDDSID
                                           KASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNID
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                                                                                      Conservative
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                                                                                                           10.3%;
                                                                                      39;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 94.5; DB 2; Length 567; Pred. No. 0.35;
                                                                                                           Score 93.5; DB 2; Pred. No. 0.66;
                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           80;
                                                                                      73;
                                                                                                                             Length
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                                                                                        Indels
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                                                                                      39;
                                                                                      Gaps
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R;Johnson, D.; Biewaid, i. gubmitted to the EMBL Data Library, July 1998 aubmitted to the EMBL Data Library, July 1998 A;Description: The sequence of C. elegans cosmid C16All. A:Reference number: Z21328
                                                                                                                                           A; Residues: 1-646 < JOH>
A; Residues: 1-646 < JOH>
A; Cross-references: UNIPROT: 076579; UNIPARC: UPI0000076553;
A; Cross-references: Strain Bristol N2; clone C16All
                                                                                                                                                                                                                                                                                                                                                                                  hypothetical protein C16A11.5 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Accession: F83128
R;Stover, C:K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, R;Stover, C:K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, R;Stover, C:K.; Folger, K.R.; Kas, adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas,
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A;Residues: 1-140 <STO>
A;Cross-references: UNIPROT:Q9HWP6; UNIPARC:UPI00000C5C2C; GB:AE004830; GB:AE004091; NIC
A;Experimental source: strain PAO1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Complete genome se
A;Reference number: A82950;
A;Accession: F83128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probable transcription regulator PA4135 [imported] - Pseudomonas aeruginosa
C;Species: Pseudomonas aeruginosa
C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 09-Jul-2004
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                                                                             A; Map position: 2
A; Introns: 35/3; 72/1;
                                                                                                                                                                                                                                                                 A; Reference number: A; Accession: T33346
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                                                                                                                                                                                                                         A;Molecule type: DNA
                                                                                                                                                                                                                                         A;Status: preliminary; translated from GB/EMBL/DDBJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 11
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                                                                                                                        Gene: CESP:C16A11.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   62 IRILRTESLQDTKPANRCCLLRH-----LLRLYLDRVFKNYQTPDHYTLRKISSLANS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          79 VQLYFADVVKNITTNDTK-----VKRLIHLYLLRFAEN---DPNLTLLSINSLQKSLS 128
l Similarity
36; Conserv
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Conservative
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                                                                                94/3; 129/3; 184/3; 205/1; 279/3; 387/3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence of Pseudomonas aeruginosa PA01, 50; MUID:20437337; PMID:10984043
                  9.0%; Score 81.5;
26.1%; Pred. No. 7.
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22;
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Pred.
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  Mismatches
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                                        BB
                                        <u>ب</u>
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  49;
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                                        Length 646;
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  Indels
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                                                                                  505/2;
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  31;
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A.; Larbig,
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  Gaps
                                                                                  540/2;
                                                                                                                                                                                       PIDN: AAC26261
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K.; Li
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protein kinase (
C;Species: Schiz
C;Date: 27-Jan-1
hypothetical protein C44Cl0.4 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change C;Accession: T19925 R;Cottage, A. submitted to the EMBL Data Library, February 1996 A;Reference number: Z19197 A;Accession: T19925 A;Status: preliminary; translated from GB/EMBL/DDBJ
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A; Molecule type: DNA
A; Residues: 1-1062 <SAU>
A; Cross references: UNITARC: UPT0000127259; EMBL: AL035537;
A; Cross references: strain 972h-; cosmid c21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: UNIPROT:P41892; UNIPARC:UPI0000127259; EMBL:X78799; NID:g521098; A;Experimental source: wildtype 972 h minus
A;Note: mRNA sequencing was also done to confirm the intron borders
A;Note: mRNA; Saunders, D.; Harris, D.; Wood, V.; Rajandream, M.A.; Barrell, B.G.
submitted to the EMBL Data Library, February 1999
A;Reference number: Z21860
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S46367
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A;Introns: 9/3; 54/3; 64/3; 86/2; 171/1; 325/1; 405/3; 427/1; 544/3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Fankhauser, C.; Simanis, V.
EMBO J. 13, 3011-3019, 1994
A;Title: The cdc7 protein kinase is a dosage dependent:
A;Reference number: S46367; MUID:94313982; PMID:8039497
A;Accession: S46367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Schizosaccharomyces pombe
C;Date: 27-Jan-1995 #sequence_revision 27-Jan-1995 #text_change 05-Oct-2004
C;Accession: S46367; T39888
                                                                                                                                                                                                             T19925
                                                                                                                                                                                                                                RESULT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            F;15-23/Region:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Description: essential for septum formation and cell division
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C; Function
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Gene: CDC7; SPDB: SPBC21.06c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-1062 < FAN>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;Keywords: ATP; cell division; phosphotransferase; protein kinase;7-259/Domain: protein kinase homology <KIN>;15-23/Region: protein kinase ATP-binding motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local (
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2-isopropylmalate synthase, probable (leuA-1) [imported] - Sulfolobus solfataricus C;Species: Sulfolobus solfataricus C;Date: 24-May-2001 #sequence_revision 24-May-2001 #text_change 09-Jul-2004 C;Accession: B90153
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A;Title: The Genome of the Natural Genetic
A;Reference number: ABZ577; MUID:21608550;
A;Accession: AD3160
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: DNA
A;Residues: 1-542 <WIL>
A;Residues: 1-542 <WIL>
A;Cross-references: UNIPROT:Q18616; UNIPARC:UPI000017B837;
A;Experimental source: clone C44Cl0
C;Genetics:
A;Gene: CESP:C44Cl0.4
A;Map position: X
A;Introns: 66/3; 229/3; 309/1; 328/3; 398/3
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A;Experimental source: strain C58 (Dupont)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hypothetical protein glnP/glnQ [imported] - Agrobacterium tumefaciens (strain
C;Species: Agrobacterium tumefaciens
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Best Local
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294, 2317-2323, 2001
rs: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan,
                                                                                                                                                                                                                                                                                                 glnP/glnQ
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322
                                                                                                                                      206 LLAGVFYLIVTVPLTHVVNAIDNRLRIGKORPSVITSGLEEV----SELDGAARASGVAF
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                                                                                                      KDGNIDIRILRRTESLQD-----TKPANRCC------LLRHL-----
                                                                                                                                                                     LLSAAFYLLWT-PSTGL----KTLNLGS---CVIATNLQEIRNGFSDIRGSVQA----
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ILIDGESILAMKPETLRRRVGMVFQHFNLFPDH----TALENVMLSLKK
                                                                     KGGSLDVRRLGMAYGDLDVLKGVDLSVKPGSVTCIIGPSGSGKSTLLRGLNRLVEPKSGD
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                                -LRLYLDRVFKNYQT-PDHYTLRKISSLANSFLTIKK 121
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                                                                                                                                                                                                        ; Score 76.5; DI
; Pred. No. 17;
23; Mismatches
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Pred. No.
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                                                                                                                                                                                                            69;
366
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R;She, Q.; Singh, R.K.; Confalonieri, F.; Zivanovic, Y.; Allard, G.; Awayez, M.J.; Chan-Jong, I.; Jeffries, A.C.; Kozera, C.J.; Medina, N.; Peng, X.; Thi-Ngoc, H.P.; Redder, F. arrett, R.A.; Ragan, M.A.; Sensen, C.W.; Van der Oost, J. submitted to GenBank, April 2001
A; Jescription: Sulfolobus solfataricus complete genome.
A;Reference number: A99139
A;Accession: B90153
A;Status; preliminary
A;Molecule type: DNA
A;Residues: 1-553 <KUR>
A;Reross-references: UNIPROT:Q980Z1; UNIPARC:UPI000006418D; GB:AE006641; NID:g13813259; F. C;Genetics:
A;Gene: leuA-1
C;Superfamily: 2-isopropylmalate synthase leuA
Search completed: December 24, 2005, 10:48:30 Job time : 39 secs
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                                                                                               402 SFDVGPASAILITLKELNI 420
                                                                                                                                                                                               352 GNDRKFVISELSGTANLVSYLQGLGIAVDKKDERL------KKALNKIKELEARGY 401
                                                                                                                                                                                                                                              104 ----YTLRKISSLAN--SFL----TIKKDLRLCHAHMTCHCGEEAMKKYSQILSHFE 150
                                                                                                                                                                                                                                                                                                                                                                                               151 KLE--PQAAVVKALGELDI 167
                                                                                                                                                                                                                                                                                                 292 QEŚLRKLREVSKIVYBILGLPPNPYQPYVGDNAFAHKAGVHVDAVMKVPRAYEHVDPSLV 351
                                                                                                                                                                                                                                                                                                                                             68 TESLQDTKFANRCC------ 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                24 GLKTLNLGSCVIATNLQEIRNGFSDIRGSVQA---KDGNID-----IRILRR
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Result
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REPARENT REP

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AD Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
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AD Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
AD Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
AD Raha S.S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
AD Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
AD Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
AD Wilting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
AD Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
AD Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
AD Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
AD AND Recompton J.E., Jones S.J.M., Marra M.A.;
AND Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;

"Generation and initial analysis of more than 15,000 full-length human
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Zkang Z., Henzel W.J.;
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Best Local S
Matches 175
            EMBL; AF224267; AAF36680.1; -; mRNA.
HSSP; O9UHDO; INIF.
Ensembl; ENSWISGO0000026416; Mus musculus.
MGI; MGI:1890473; I120.
GO; GO:0005615; C:extracellular space; TAS.
GO; GO:0045517; F:interleukin-20 receptor bindis
GO; GO:0045617; F:hemopoiesis; NAS.
GO; GO:0045618; P:positive regulation of epiders
GO; GO:0045618; P:positive regulation of keratis
GO; GO:0045618; P:positive regulation of tyrosis
GO; GO:0045618; P:positive regulation of tyrosis
GO; GO:0045618; D:positive regulation of tyrosis
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CONFLICT
SEQUENCE
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28-FEB-2003
10-MAY-2005
                                                                                                                                                                                                                                                                                                                                                                                                         epidermal function.";
Cell 104:9-19 (2001).
-i- FUNCTION: Cytokine that
                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=21097717; PubMed=11163236; DOI=10.1016/S0092-8674(01)00187-8; Blumberg H., Conklin D., Xu W.F., Grossmann A., Brender T., Carollo S., Eagan M., Foster D., Haldeman B.A., Hammond A., Haugen H. Jelinek L., Kelly J.D., Madden K., Maurer M.F., Parrish-Novak J., Frunkard D., Sexson S., Sprecher C., Waggie K., West J., Whitmore T.E., Yao L., Kuechle M.K., Dale B.A., Chandrasekher Y.A.; "Interleukin 20; discovery, receptor identification, and role in epidermal function.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IL20 MOI
Q9JKV9;
                                                                                                                                                                                                                                                                               This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no restruse as long as its content is in no way modified and this s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Name=1120; Synonyms=Zcyto10;
Mus musculus (Mouse).
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28-FEB-2003 (Rel. 41, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Interleukin-20 precursor (IL-20) (Four alpha helix cytokine ZCYTO10).
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SUBCELLULAR LOCATION: Secreted.
SIMILARITY: Belongs to the IL-10
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No. 1.8e-80;
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                                                                                of epidermal cell diffe. of keratinocyte differe. of tyrosine phosphoryla.
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lires; Rodentia;
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Matches 131
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28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
10-MAY-2005 (Rel. 47, Last annotation update)
Interleukin-19 precursor (IL-19) (Melanoma differentiation protein-like protein) (NG.1).
Name=IL19; Synonyms=ZNDA1;
Homo sapiens (Human)
                                                                                                 Kotenko S.
Submitted
                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE.

MEDLINE=22257669; PubMed=12370360;

Liao Y.-C., Liang W.G., Chen F.W., Hsu

"IL-19 induces production of IL-6 and
apoptosis through TNF-alpha.";
J. Immunol. 169:4288-4297(2002).
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PROSITE; I
Cytokine;
SIGNAL
CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.
MEDILINE=21040165; PubMed=11196675; DOI=10.1038/sj.gene.6363714;
MEDILINE=21040165; PubMed=11196675, DOI=10.1038/sj.gene.6363714;
Milagher G., Dickensheets H., Eskdale J., Izotova L.S.,
Mirochnitchenko O.V., Peat J.D., Vasquez N., Pestka S., Donnell
                    NUCLEOTIDE SEQUENCE [GENOMIC DNA], Rieder M.J., Carrington D.P., Chung Nickerson D.A.;
                                                                                                                                                                                                                    Sexson S.,
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                                                                                   "Ηοπο
                                                                                                                                                                                                                                                                                                                                                                                                                                  "Cloning, expression and initial characterization (IL-19), a novel homolog of human interleukin-10 of the Immun. 1:442-450(2000).
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n S., Smith D., Lok S.,
sapiens homolog of mel
tted (OCT-1999) to the
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                                                                                                 .V., Pestka
(JUN-2001)
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                                                                                                                                      SEQUENCE.
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      NHLBI
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176
126
132
    HL66682
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75.3%;
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S., Pownder T., O'Hara P.
melanoma differentiation
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Mismatches 2
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                                    AND VARIANT
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TNF-alpha and
    genomic
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applications,
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MKLQCVSLWLLGTILILCSVDNHGLR----

MKASSLAFSILSAAFYILWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNI

RCLISTDMHHIEESFQEIKRAIQAKDTFP

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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                            GO; GO:0005576; C:extracellular region; N
GO; GO:0005125; F:cytokine activity; TAS:
GO; GO:0006955; P:immune response; NAS.
GO; GO:0007165; P:signal transduction; NA
InterPro; IPR012351; Cytokine 4 hlx.
InterPro; IPR000098; Interleukin 10.
Pfam; PF00726; II10; 1.
SMAXT; SM00188; II10; 1.
                                                                                                                                                                                                                                                                                                     Glycoprotein;
SIGNAL
CHAIN 2
                                                                                                                                                 CARBOHYD
CARBOHYD
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J. Biol. Chem. 278:3308-3313(2003).
-!- FUNCTION: May play some important roles
Up-regulates IL-6 and TWF-alpha and indusimilarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This Swiss-Prot entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
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PubMed=12403790; DOI=10.1074/jbc.M208602200;
Chang C., Magracheva E., Kozlov S., Fong S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PubMed=15340161; DOI=10.1110/ps.04682504;
Zhang Z., Henzel W.J.;
"Signal peptide prediction based on analy:
verified cleavage sites.";
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Human chromosome 1 i
Submitted (FEB-2000)
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Ensembl; ENSG00000142224; H
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                                                                                                                                                                                                                                                                                                                                                                               3D-structure;
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AF453946; AAN49906.1; -; mRNA.

AF192498; AAF06663.1; -; mRNA.

AY040367; AAK91776.1; ALT_INIT; mRNA.

AF390905; AAK64498.1; -; Genomic_DNA.

AL049615; CAB72342.1; -; Genomic_DNA.
       69;
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N-linked (GlcNAc
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13-SEP-2005 (TrEMBLrel. 31, Last sequence up
13-SEP-2005 (TrEMBLrel. 31, Last annotation
13-SEP-2005 (TrEMBLrel. 31, Last annotation
Interleukin 10 family protein.
Brachydanio rerio (Zebrafish) (Danio rerio)
                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eu
Actinopterygii; Neopterygii; Teleostei; Ostariophysi;
Cyprinidae; Danio.
                                                                                                                                                                                                                                    Q4LDR4 BRARE PRELIMINARY;
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Pfam; PF00726; IL10; 1.
PROSITE; PS00520; INTERLEUKIN_10; UNKNOWN_1.
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EMBL; AL513315; CAH71814.1; -; Genomic_DNA.
SMR; QSVUT3; 60-215.
Ingembl: ENGCOCOLOR
                                                         NCBI_TaxID=7955;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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         NUCLEOTIDE SEQUENCE Ram S., Daisuke I.,
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BRARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       embl; ENSG00000142224; Homo sapiens.
GG:0005615; C:extracellular space; IEA.
GO:0005125; F:cytokine activity; IEA.
GO:0006955; P:immune response; IEA.
                                                                                                                                                                                                                                                                                                                                                                                   121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        69;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DIRILRRTESLODTKFANRCCLLRHLLRLYLDRVFKNYQTFDHYTLRKISSLANSFLTIK 120
                                                                                                                                                                                                                                                                                                                                                                                                                NTILSTLETLQIIKPLDVCCVTKNLLAFYVDRVFKDHQEPNPKILRKISSIANSFLYMQ
                                                                                                                                                                                                                                                                                                                                                                                                                                           MKASSLAFSLLSAAFYLLWTPSTGLKTLNLGSCVIATNLQEIRNGFSDIRGSVQAKDGNI
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                                                                                                                                                                                                                                                                                                                                                                                   KDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMEE 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MKLQCVSLWLLGTILILCSVDNHGLR-----RCLISTDMHHIEESFQEIKRAIQAKDTFP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  215 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24567 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chordata;
         Sakai M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    %; Score 337.5;
%; Pred. No. 1.2e
36; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Last sequence up
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Created)
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oglires; Primates; Cata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  D79D6B0F11DA74E4 CRC64;
                                                                                                                                                                                                                                                       PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ::
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                       Euteleostomi;
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                                                                                                    Cypriniformes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             215;
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IL19_MC
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Best Local S
Matches 67
         CHAIN
CARBOHYD
CARBOHYD
                                                               Apoptosis;
SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOUSE
                                                                                                                                 MGI; MGI:1894,4; 1...

GO; GO:0006917; P:induction of apoptos

GO; GO:0042226; P:interleukin-6 biosyn

GO; GO:0006800; P:oxygen and reactive

InterPro; IPR012351; Cytokine 4 hix.

InterPro; IPR000098; Interleukin_10; 1.
                                                                                                                                                                                             Ensembl; ENSMUSG00000016524; Mus MGI; MGI:1890472; Il19.
GO; GO:0006917; P:induction of ap GO; GO:0042226; P:interleukin-6 b GO; GO:0006800; P:oxygen and reac GO; GO:0006800; P:oxygen and reac
                                                                                                                                                                                                                                                                                        EMBL; AF453945; AAN40905.1;
HSSP; Q9UHD0; 1N1F.
SMR; Q8CJ70; 23-170.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN-Swiss Webster / NIH;
MEDLINE-22257669; PubMed-12370360;
Liao Y.-C., Liang W.G., Chen F.W., Hsu
"II-19 induces production of II-6 and
apoptosis through TNF-alpha.";
J. Immunol 169:4288-4297(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IL19_MOI
Q8CJ70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Evolution of IL-10 family genes in teleosts."; Submitted (JUL-2005) to the EMBL/GenBank/DDBJ database EMBL; AB219579; BAED60181.1; -; mRNA.
SEQUENCE 168 AA; 19110 MW; 807D18AD736C709B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                   This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no restude as long as its content is in no way modified and this use as long as its content is in no way modified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-MAY-2005 (Rel. 47, Created)
10-MAY-2005 (Rel. 47, Last sequence up
10-MAY-2005 (Rel. 47, Last annotation
                                                                                                                           SMART;

    -i- FUNCTION: May play some important roles
        Up-regulates IL-6 and TNF-alpha and indu
-i- SUBCELULAR LOCATION: Secreted.
    -i- SIMILARITY: Belongs to the IL-10 family.

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Muroidea; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=Il19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Interleukin-19
                                                                                                        PROSITE; PS00520;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                67;
                                                                                                                         SM00188;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENTRIQIKSIQTAYEKIDQAAGTVKAIGELDSLLEWIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWME 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ---SCCFLSQLLHFYMDTVFISYTSSHSLHRRTTSVLANSFLSISKDLRVCHANAHCECG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PANRCCLLRHLLRLYLDRVFKNYQTFDHYTLRKISSLANSFLTIKKDLRLCHAHMTCHCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LWDAAQG-RRLHLGSCKVNIHTHELRHHFQYVRQGMISGDDHKGIRLLRKDVMSSLQATE
                                                                                   Cytokine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE,
             1
25
56
127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  precursor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD;
                                                                                                                         IL10;
                                                                                 INTERLEUKIN 10; FALSE NEG. ne; Glycoprotein; Signal.
             24
176
56
127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Glires; Rodentia; Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                33.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AND
                                                                 Glycoprotein; Sic
24 Potentia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (IL-19).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FUNCTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27;
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Pred. No. 1.1e
27; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus
                                                                                                                                                                                                                  of apoptosis; IDA.
in-6 biosynthesis;
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         Interleukin-19.
N-linked (GlcNAc.
N-linked (GlcNAc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
                                                                                                                                                                                                                                                                                                                                mRNA
                                                                                                                                                                                                                                                                        musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Heu
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No. 1.1e-21;
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TNF-alpha and
                                                                                                                                                                                                   oxygen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    induces
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  apoptosis
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             (Potential). (Potential).
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this statement
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                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL
                                                                                                                                                                                                                                                                                                                                                                                      statement
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                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Comparative genomic analysis reveals independent expansion of Ilineage-specific gene family in vertebrates: The class II cytok I receptors and their ligands in mammals and fish.";

BMC Genomics 4:29-29(2003).

EMBL; AY294557; AAP57414.1; -; Genomic_DNA.

REMBL; AY294558; AAP57416.1; -; mRNA.

GO; GO:0005615; C:extracellular space; IEA.

GO; GO:0005615; C:extracellular space; IEA.

GO; GO:0005125; F:cytokine activity; IEA.

GO; GO:0005125; F:cytokine activity; IEA.

R GO; GO:0005125; F:cytokine activity; IEA.

R InterPro; IPR000098; Interleukin_10.

Pramm: PR00726; IL10; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                           ProDom; PD003687; Interleukin_10; 1.
SMART; SM00188; IL10; 1.
SEQUENCE 175 AA; 19838 MW; DE6FA
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Mogensen K., Monneron D.;
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Pred. No. 8.4e-18;
3; Mismatches 53;
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rpha; Tetraodontiformes;
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DT 01-DEC-2
DT 01-MAR-2
DE Melanoma
GN Name=112
OS Musme=12
OC Eukaryot
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Matches 56
  NUCLEOTIDE SEQUENCE.
STRAIN=129/SvJ;
Madireddi M.T., Lin J
Submitted (FEB-2000)
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SEQUENCE
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01-DEC-2001 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
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Q925S4;
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GG; GO:0005615; C:extracellular space;
InterPro; IPR000098; Interleukin 10; 1.
ProDom; PD003687; Interleukin 10; 1.
SMART; SM00188; IL10; 1.
PROSITE; PS00520; INTERLEUKIN 10; 1.
                                                                                                                                                              Eukaryota; Metazoa; Ch
Mammalia; Eutheria; Eu
Muridae; Murinae; Mus.
                                                                                                                                                                                                                                              Mus musculus (Mouse)
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Mammalia; Eutheria; Eu
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Ensembl; ENSMUSG00000026420; Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Th2-specific cytokine Name=Il24;
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01-MAR-2004
                                                                                                                                  NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                          Name=I124;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=21240641; PubMed=11342597;
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     J.,
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                                                                                                                                                                                                                     Chordata;
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FISP.
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19,
26,
     Su 2
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26, Last annotation update)
associated gene-7.
     Z., Shay J.W., Huberman E.,
EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                           Created)
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Pred. No. 2.6e
30; Mismatches
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                                                                                                                                                                                         Sciurognathi;
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RESULT 10
IL24_HUMAN
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Best Loc
Matches
                                                                                                 wenomic structure, chromosomal localization and expressio melanoma differentiation associated gene-7 (mda-7): potent relationship with cellular senescence.";
Submitted (FEB-2000) to the EMRT /~-
[4]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q13077; Q96DB0; Q96KG4;
01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
10-MAY-2005 (Rel. 47, Last annotation update)
11-Rerleukin-24 precursor (Suppression of tumorigenicity
(Melanoma differentiation-associated protein 7) (MDA-7).
Name=IL24; Synonyms=MDA7, ST16;
Rieder M.J., Armel I.A., VI. O., Nickerson D.A.;
Poel C.L., Toch B.J., Yi O., Nickerson D.A.;
Poel C.L., HIBI HL66682 program for genomic applications,
"SeattleSNPs. NHIBI HL66682 program for genomic applications,
"SeattleSNPs. WA (URL: http://pga.gs.washington.edu).";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -|- SUBCELLULAR LOCATION: Secreted (By similarity).
EMBL; AF235006; AAK52590.1; -; mRNA.
Ensembl; ENSMUSG0000002641; -ws musculus.
MGI; MGI:2135548; I124.
GG; GO:0005615; C:extracellular space; TAS.
                                                                                                                                                                                                                                                                                                      MEDLINE=96132699; PubMed=8545104;
Jiang H., Lin J.J., Su Z.-Z., Goldstein N.I., Fisher P.B.;
"Subtraction hybridization tiffies a novel melanoma differentiation
associated gene, mda-7, modulated during human melanoma
differentiation, growth and progression.";
Oncogene 11:2477-2486(1995).
                                                                                                                                                                                                                                 Peat J., Kube D., Eskdale J., Jueliger "The human MDA-7 gene.";
Submitted (JUN-2000) to the EMBL/GenBan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR000098; Interleukin | ProDom; PD003687; Interleukin_10; SMART; SM00188; IL10; 1.
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                                                                                           NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                           TISSUE=Melanoma;
                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE; PS00520; INTERLEUKIN_10; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
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                                                                                         SEQUENCE
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                                               Armel T.Z., v
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20812 MW; 05CA43872D53D555
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Homin
                                                                                         [GENOMIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24.2%;
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Pred. No. 3
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                                                                                           AND
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                                                                                           VARIANTS
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les 71;
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                                                                                                                                                                                                                                                              Gallagher
                                                                                                                                                n and expression o (mda-7): potential
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                                                                                                                                                                             Fisher P.B.;
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A Strausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., A Klausner R.D., Collins F.S., Beetow K.H., Shenmen C.M., Schuler G.D., A Altschul S.F., Zeeberg B., Buetow K.H., Shenmen C.M., Schuler G.D., A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., A Grant G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Stapleton M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C., A Stapleton M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C., A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Raha S.S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H., Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A., Rah Hitting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., A Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Schein J.E., Jones S.J.M., Marra M.A.; "Generation and initial analysis of more than 15,000 full-length human and generation and initial analysis of more than 15,000 full-length human and generation and malysis of more than 15,000 full-length human and generation and malysis of more than 15,000 full-length human
                                                                       InterPro; IPR012351; Cytokine 4 hlx. InterPro; IPR012351; Cytokine 4 hlx. InterPro; IPR000098; Interleukin 10. ProDom; PD003687; Interleukin 10; 1. SMART; SM00188; IL10; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE-9904566; PubMed-9826712; DOI=10.1073/pnas.95.24.14400; Su Z.-Z., Madireddi M.T., Lin J.J., Young C.S.H., Kitada S., Reed J.C., Goldstein N.I., Fisher P.B.; "The cancer growth suppressor gene mda-7 selectively induces apoptosis in human breast cancer cells and inhibits tumor growth in nude mice."; proc. Natl. Acad. Sci. U.S.A. 95:14400-14405(1998).

-i- FUNCTION: Has antiproliferative properties in human melanoma cells and may contribute to terminal cell differentiation. May also function as a negative regulator of melanoma progression. Formation in vivo in nude mice. They found that it selectively induces apoptosis in human breast cancer cells.

-i- SUBCELLULAR LOCATION: Secreted (Potential).

-i- TISSUE SPECIFICITY: Up-regulated in melonoma cells induced to
                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; U16261; AAA91780.1; -; mRNA.

EMBL; AF276916; AAG41401.1; -; Genomic_DNA.

EMBL; AF235005; AAK52589.1; -; Genomic_DNA.

EMBL; AY062931; AAK34146.1; -; Genomic_DNA.

EMBL; BC009681; AAH09681.1; -; mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This Swiss-Prot entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict use as long as its content is in no way modified and this statuse as long as its content is in no way modified and this statuse as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as long as lon
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PubMed=15340161; DOI=10.1110/ps.04682504;
Zhang Z., Henzel W.J.;
"Signal peptide prediction based on analy
verified cleavage sites.";
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                                                                                                                                                                                                                                                                                                                                       Ensembl; ENSG00000162892; HGNC; HGNC:11346; IL24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and mouse cDNA sequences."
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                                                            PROSITE; PS00520; INTERLEUKIN_10;
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                                Cytokine;
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Feingold E.A., Gro
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Grouse L.H., Derge J.G.,
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CARBOHYD
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                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
Kalnine N., Chen X., Rolfe
Koundinya M., Raphael J.,
Phelan M., Farmer A.;
                                                                                                                                                                                                                                                                                                                                NCBI
                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                    Submitted (MAY-2003) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                    Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
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                                                                                                                                                                     Cytokine
                                                                                                                                                                              GO; GO:0006955;
                                                                                                                                                                                                                                                         Phelan M., Farmer A.; "Cloning of human full-length CDSs in BD
                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human)
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                                                                                                                                                                          SUBCELLULAR LOCATION: Secreted (By 8 L; BT007156; AAP35820.1; -; mRNA. GO:0005576; C:extracellular region; GO:0005125; F:cytokine activity; IER GO:0006955; P:immune response; IEA.
                                                                                                                                                                                                                                                                                                                               TaxID=9606;
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 KDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVKALGELDILLQWMEE 174
                     RLLQQ-EVLQNVSDAESCYLVHTLLEFYLKTVFKNYHNRTVEVRTLKSFSTLANNFVLIV
                                          RILRRTESLODTKPANRCCLLRHLLRLYLDRVFKNY--QTPDHYTLRKISSLANSFLTIK
                                                                 LGFTL-----LLWSQVSGAQGQEFHFGPCQVKGVVPQKLWEAFWAVKDTMQAQDNITSA
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126
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23825
                                                                                                                      23.8%;
                                                                                                                                                        23912 MW;
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l J., Moreira D., Kelley T., LaBaer J., Lin
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32.8%;
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N-linked (GlcNAG. . .) (Pote N-linked (GlcNAG. . .) (Pote N-linked (GlcNAG. . .) (Pote Y-> H (in dbSNP:1150258).

/FTId=VAR 011974.

H -> R (in dbSNP:3093431).
/FTId=VAR 013097.

V -> L (in dbSNP:3093446).
/FTId=VAR 013098.
/FTId=VAR 013098.
                                                                                                                                                                                                                                                                                                                                                                                               Last sequence update)
Last annotation update
                                                                                                                                                                                                                                                                                                                                                                                                                      Created)
                                                                                                                       Score 217;
Pred. No. 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 217; DB 1;
Pred. No. 7.3e-13
                                                                                                                                                        17687A467B9CB551 CRC64;
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SEQUENCE
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"Identification of a novel ligand-receptor pair constitutively
activated by ras oncogenes.";
J. Biol. Chem. 275:24436-24443(2000).
-!- SUBCELLULAR LOCATION: Secreted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ensembl; ENSRNOG00000004470; Rattus norvegicus. GO; GO:0048144; P:fibroblast proliferation; TAS. GO; GO:0042060; P:wound healing; TAS. InterPro; IPR012351; Cytokine 4 hlx. InterPro; IPR000098; Interleukin_10. Probom; PD003687; Interleukin_10; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This Swiss-Prot entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
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28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Cytokine-like protein Mob-5 precursor.
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna
Muroidea; Muridae; Murinae; Rattus.
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MEDLINE=20387329; PubMed=10825166; DOI=10.1074/jbc.M001958200;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=10116;
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1 28 Po
29 183 Cy
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21096 MW;
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Pred. No. 2.2e-10;
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N-linked (GlcNAc. . .) (Pote;
AF7A11466C491AC4 CRC64;
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12, 12, 26,

Created)
Last sequence update)
Last annotation update)

PRELIMINARY,

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MEDLINE=99308986; PubMed=10381256;

MEDLINE=99308986; PubMed=10381256;

DOI=10.1002/(SICI)1097-4644(19990701)74:1<1::AID-JCB1>3.3.CO;2-D;

SOO C., Shaw W.W., Freymiller E., Longaker M.T., Bertolami C.N.,

Chiu R., Tieu A., Ting K.;

Chiu R., Tieu A., Ting K.;
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C49a.

Rattus norvegicus (Rat).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
                                                                                                    Immune Response to Eimeria maxima.";

J. Immunol. 173:2675-2682(2004).

EMBL; AJ621254; CAF18432.1; -; mRNA.

GO; GO:0005576; C:extracellular region;

GO; GO:0005125; F:cytokine activity; IE;

GO; GO:0006955; P:immune response; IEA.
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25-OCT-2004 (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -!- SUBCELLULAR LOCATION: Secreted (By similarity).
EMBL, AF004774; AAB691/11.1; -; mRNA.
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:0005125; F:cytckine activity; IEA.
GO; GO:0006955; P:immune response; IEA.
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J. Cell. Biochem. 74:1-10(1999).
                               InterPro; IPR012351; Cytokine 4 hlx.
InterPro; IPR012352; IL-10 add helix.
InterPro; IPR000098; Interleukin_10.
                                                                                                                                                                                                                                                                                     Rothwell L., Young J., Archer A., Smith A.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q6A2H5 CHICK PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9031;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Archosauria; Aves;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gallus gallus (Chicken)
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47; Conserv
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                                                                                                                                                                                                                                                                                                       J.,
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J. Immunol. 173:2675-2682(2004).

EMBL; AJ621614; CAF21727.1; -; Genomic DNA.

GO; GO:0005576; C:extracellular region; IEA.

GO; GO:0005525; F:cytckine activity; IEA.

GO; GO:0006955; P:immune response; IEA.

InterPro; IPR012351; Cytckine 4 hlx.

InterPro; IPR012352; IL-10 add helix.

InterPro; IPR00098; InterTeukIn_10.

Pfam; PF00726; IL10; 1.
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Q6A2H4;
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ProDom; PD003687; Interleukin_
SMART; SM00188; IL10; 1.
                                                                                                                                                                                                                                                                                                                                                                                       PRINTS; PR01294; INTRLEUKIN10.
ProDom; PD003687; Interleukin_10; 1.
SMART; SM00188; IL10; 1.
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Eukaryota; Metazoa; Chordata;
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Archer A., Smith A.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Archosauria; Aves; Neognathae;
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117
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; Galliformes; Phasianidae; Phasianinae;
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7e-09;
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ALIGNMENTS

1 (bases 1 to 926)
Conklin, D.C., Haldeman, B.A. and Grossmann, A. Mammalian cytokine-like polypeptide-10
Patent: JP 2001524313-A 1 04-DEC-2001;
ZYMOGENETICS INC
OS Homo sapiens (human)
PN JP 2001524313-A/1
PD 04-DEC-2001
PD 04-DEC-2001
PF 25-NOV-1998 JP 2000522245
PF 26-NOV-1997 US 08/979156
PR 26-NOV-1997 US 08/979156
PR 26-NOV-1997 US 08/979156
PR 26-NOV-1997 US 08/979156
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PR 26-NOV-1997 US 08/979156
PR 26-NOV-1997 US 08/979156
PR 26-NOV-1997 US 08/979156
PR 26-NOV-1997 US 08/979156 Homo sapiens (human) BD087297.1 GI:22632907 JP 2001524313-A/1. BD087297 926 bp DNA Mammalian cytokine-like polypeptide-10. Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; łominidae; Homo. 25-NOV-1998 JP 2000522245 26-NOV-1997 US 08/979156 DARRELL C CONKLIN.BETTY A HALDEMAN,ANGELIKA GROSSMANN PC 15/09,A01K67/027,A61K38/00,A61K48/00,A61P1/02,A61P11/00, I C12N15/00 A61P35/00,A61P37/02,A61P43/00,C07K14/52,C07K16/24,C07K16/42 linear PAT 27-AUG-2002 Š

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                                                                                             Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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                               Conklin, D.C., Haldeman, B.A. and Grossmann, Mammalian cytokine-like polypeptide-10 Patent: Ep 142433-A 1 02-JUN-2004; Zymogenetics Inc (US)
                                                                                        Hominidae; Homo
                                                                                                                                       Homo
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Mammalian
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Sequence 1
CS109174
CS109174.1
              Moore, M.D.
Anti-il-20 receptor antibodies
                                         Xu, W., Kindsvogel, W.R., Ch
Dillon, S.R., Lehner, J.M.,
                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Euarchontoglires; Primates; Cata
using in inflammation
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                                           Chen, Z., Hughes, S.D., Chandrasekher, Y.A., Siadak, A.W., Sivakumar, P.V. and
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
Hominidae; Homo.
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/brottein_id="A2J09906.1"
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1 (bases 1 to 926)

Blumberg, H., Conklin, D., Xu, W.F., Grossmann, A., Brender, T.,
Blumberg, Eagan, M., Poster, D., Haldeman, B.A., Hammond, A.,
Carollo, S., Eagan, M., Poster, D., Madden, K., Maurer, M.F.,
Haugen, H., Jelinek, L., Kelly, J.D., Madden, K., Maurer, M.F.,
Parrish-Novak, J., Prunkard, D., Sexson, S., Sprecher, C., Waggie, K.,
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Conklin,D.C. and Haldeman,B.A.
Conklin,D.C. and Haldeman,B.A.
Mammalian cytokine-like polypeptide-10
Patent: US 6576743-A 1 10-JUN-2003,
ZymoGenetics, Inc.; Seattle, WA
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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Chandrasekher,Y.A.
Interleukin 20: discovery, recep
epidermal function
cell 104 (1), 9-19 (2001)
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Direct Submission
Submitted (12-JAN-2000) Biomolecular Informatics,
Inc., 1201 Eastlake Ave. East, Seattle, Wa. 98102.
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/allele="1"
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Sequence 18 from Patent
CS123418
CS123418.1 GI:70911957
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Location/Qualifiers
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                                             GluGluThrGlu 144
                                                                                                GluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet 140
                                                                                                                                                      HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu 120
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                                                                                                                                                                                                                                                                                                                                                                                                                              CysVallleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer
                                                                                                                                      CACTGTGGTGAGGAAGCAATGAAAAAAATACAGCCAGATTCTGAGCCACTTCGAAAAACTG
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                                                                                                                                                                                                                                                                ArgValPheLysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIleSerSerLeu
                                                                                                                                                                                                                                                                                                                           AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp 60
                             GAGGAGACCGAA 459
                                                                                                                                                                                           GCCAACTCCTTCCTGACCATCAAAAAAAGACCTGCGTCTGTGTCACGCCCACATGACCTGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
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1 (bases 1 to 531)
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Strausberg, R. D., Feingold, E. A., Grouse, L. H., Derge, J. G.,

Klausner, R. D., Collins, F. S., Wagner, L., Schaefer, C. F., Bhat, N. K.,

Altschul, S. F., Zeeberg, B., Buetow, K. H., Schaefer, C. F., Bhat, N. K.,

Hopkins, R. F., Jordan, H., Moore, T., Max, S. I., Wang, J., Hsheh, F.,

Diatchenko, L., Marusina, K., Farmer, A. A., Rubin, G. M., Hong, L.,

Scheetz, T. E., Brownstein, M. J., Usdin, T. B., Toshiyuki, S.,

Carninci, P., Prange, C., Raha, S. S., Loquellano, N. A., Peters, G. J.,

Abramson, R. D., Mullahy, S. J., Bosak, S. A., McEwan, P. J.,

McKernan, K. J., Malek, J. A., Gunaratne, P. H., Richards, S.,

Worley, K. C., Hale, S., Garcia, A. M., Gay, L. J., Hulyk, S. W.,

Villalon, D. K., Muzny, D. M., Sodergren, B. J., Lu, X., Gibbs, R. A.,

Fahey, J., Helton, E., Ketteman, M., Young, A. C., Shevchenko, Y.,

Bouffard, G. G., Blakesley, R. W., Touchman, J. W., Green, E. D.,

Bouffard, G. G., Blakesley, R. W., Touchman, J. W., Green, E. D.,

Butterfield, Y. S., Krzywinski, M. I., Skalska, U., Smailus, D. B.,

Schnerch, A., Schein, J. E., Jones, S. J. and Marra, M. A.

Mammallan Gene Collection Program Team

Generation and initial analysis of more than 15,000 full-length

human and mouse cDNA sequences

Proc. Natl. Acad. Sci. U.S. A. 99 (26), 16899-16903 (2002)
                                                                                                                                                                                                              Web site: http://www.hgsc.bcm.tmc.edu/cdna/
Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Loulseged, H.,
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny, D.M., Nanavati,
A.N., Gibbs, R.A.
Clone distribution: MGC clone distribution information can be for through the I.M.A.G.B. Consortium/LINI at: http://image.llnl.gov Series: IRBR Plate: 1 Row: c Column: 2.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Direct Submission
Submitted (29-APR-2004) National Institutes of Health, Mammalian Gene Collection (MGC), Bethesda, MD 20892-2590, USA
NIH-MGC Project URL: http://mgc.nci.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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Hominidae; Homo.

1 (bases 1 to 531)

Strausberg,R.L., Feingold,E.A., Grouse,L.H.,
Klausner,R.D., Collins,F.S., Wagner,L., Shenn
                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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Hytlrkisslansfltikkdlrlchahwtchcgeeamkkysqilshfeklepqaavvk
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L.H., Derge,J.G.,
Shenmen,C.M., Schuler,G.D.,
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Altschul, S. F., Zeeberg, B., Buetow, K. H., Schaefer, C. F., Bhat, N. K., Hopkins, R. F., Jordan, H., Moore, T., Max, S. I., Wang, J., Hsieh, F., Diatchenko, L., Marusina, K., Faamer, A. A., Rubin, G. M., Hong, L., Stapleton, M., Soares, M. B., Bonaldo, M. F., Casavant, T. L., Scheetz, T. E., Brownstein, M. J., Usdin, T. B., Toshiyuki, S., Carninci, P., Prange, C., Raha, S. S., Loquellano, N. A., Peters, G. J., Abramson, R. D., Mullahy, S. J., Bosak, S. A., McEwan, P. J., Abramson, R. J., Mullahy, S. J., Bosak, S. A., McEwan, P. J., McKernan, K. J., Malek, J. A., Gunaratne, P. H., Richards, S. M., Worley, K. C., Hale, S., Garcia, A. M., Gay, L. J., Hulyk, S. M., Villalon, D. K., Muzny, D. M., Sodergren, E. J., Lu, X., Gibbs, R. A., Fahey, J., Helton, E., Ketteman, M., Madan, A., Rodrigues, S., Sanchez, A., Whiting, M., Madan, A., Young, A. C., Shevchenko, Y., Bouffard, G. G., Blakesley, R. W., Touchman, J. W., Green, E. D., Dickson, M. C., Rodriguez, A. C., Grimwood, J., Schmutz, J., Myers, R. M., Butterfield, Y. S., Krzywinski, M. I., Skalska, U., Smailus, D. E., Schnerch, A., Schein, J. E., Jones, S. J. and Marra, M. A. Generation and initial analysis of more than 15,000 full-length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Baylor Human Genome Sequencing Center
Tissue Procurement: Baylor Human Genome Sequencing (
cDNA Library Preparation: Baylor Human Genome Sequencing (
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Baylor College of Medicine Human Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Clone distribution: MGC clone distribution information can be for through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Series: IRBR Plate: 1 Row: d Column: 2.

Location/Qualifiers
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Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Huly
Kowis, C.R., Sneed, A.J., Martin, R.G., Muz
A.N., Gibbs, R.A.
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Submitted (29-APR-2004) National Institutes of Health, Mammalian Gene Collection (RGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
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Contact: MGC help desk
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Nanavati,
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RS Strausberg, L., Feingold, E.A., Grouse, L.H., Derge, J.G.,

Klausner, R.D., Collins, F.S., Wagner, L., Shenmen, C.M., Schuler, G.D.,

Altschul, S.F., Zeeberg, B., Buctow, K.H., Schaefer, C.F., Bhat, N.K.,

Hopkins, R.F., Jordan, H., Moore, T., Max, S.I., Wang, J., Hsieh, F.,

Diatchenko, L., Marusina, K., Farmer, A.A., Rubin, G.M., Hong, L.,

Stapleton, M., Soares, M.B., Bonaldo, M.F., Casavant, T.L.,

Scheetz, T.E., Brownstein, M.J., Usdin, T.B., Toshiyuki, S.,

Carninci, P., Prange, C., Raha, S.S., Loquellano, M.A., Peters, G.J.,

Abramson, R.D., Mullahy, S.J., Gunaratne, P.H., Richards, S.,

McKernan, K.J., Malek, J.A., Gunaratne, P.H., Richards, S.,

Worley, K.C., Hale, S., Garcia, A.M., GayL, J., Hilyk, S.W.,

Villalon, D.K., Muzny, D.M., Sodergren, E.J., Lu, X., Gibbs, R.A.,

Fahey, J., Helton, E., Ketteman, M., Madan, A., Rodrigues, S.,

Sanchez, A., Whiting, M., Madan, A., Young, A.C., Shevchenko, Y.,

Bouffard, G.G., Blakesley, R.W., Touchman, J.W., Green, E.D.,

Dickson, M.C., Rodriguez, A.C., Grimwood, J., Schmutz, J., Myers, R.M.,

Butterfield, Y.S., Krzywinski, M.I., Skalska, U., Smailus, D.E.,
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Series: IRBR Plate: 1 Row: e Column: 2.
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Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Loulseged, H.,
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny, D.M., Nanavati,
A.N., Gibbs, R.A.
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Proc. Natl. Acad. Sci. U.S.A. 99 (26), 16899-16903 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Center code: BCM-HGSC
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                                                                                                                                                                                                                                                                                                                            /gene="IL20"
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/product="interleukin 20, p
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HYTLRKISSLANSFLTIKKDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVK
                                                                                                                                                                                                                                         ALGELDILLQWMBETE"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="synonyms: ZCYTO10,
/db_xref="GeneID:50604"
/db_xref="MIM:605619"
1. .531
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/tlssue_type="PCR rescued clones"
/clone_lib="WIH_MGC_244"
/note="Vector: pPCR-Script Amp_SK(+)"
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db_xref="taxon:9606"
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9**7** 21

ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln 40

CysValIleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer 20

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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Baylor Human Genome Sequencing Center
cDNA Library Preparation: Baylor Human Genome Sequencing Center
                                                                                   Direct Submission
Submitted (29-APR-2004) National Institutes of Health, Mammalian
Submitted (MGC), Cancer Genomics Office, National Cancer
Gene Collection (MGC), Cancer Genomics Office, National Cancer
Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                           NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
                                                                                                                                            Director MGC Project.
                                                                                                                                                                                      human and mouse cDNA sequences
Proc. Natl. Acad. Sci. U.S.A.
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny,
A.N., Gibbs, R.A.
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HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
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/mol_type="mRNA"
/db_xref="taxon:9606"
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/fissue_type="PCR rescued clones"
/clone_lib="NNH_MGC_244"
/note="Vector: pPCR-Script Amp_SK
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/codon_start=1
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IM Mammalian Gene Collection Program Team
Generation and initial analysis of more than 15,000 full-length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Director MGC Project.

Direct Submission

Submitted (29-APR-2004) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Gene Collection (MGC), Cancer Genomics Office, MD 20892-2590, Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                           Web site: http://www.hgsc.bcm.tmc.edu/cdna/
Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., J
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny, D.M.,
A.N., Gibbs, R.A.
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                                                                                                                                                                                                                                                                                                                                                          Tissue Procurement: Baylor Human Genome Sequencing
                                                                                                                                                                                                                                                                                                                                                                                               NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human and mouse cDNA sequences Proc. Natl. Acad. Sci. U.S.A.
                                                                                                                                                                                                                                                            Center code: BCM-HGSC
                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
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Mammalia; Eutheria;
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IMAGE:7262164), complete
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                                                                             distribution: MGC clone distribution gh the I.M.A.G.E. Consortium/LLNL at: s: IRBR Plate: 1 Row: g Column: 2.
/organism="Homo sapiens"
/mol_type="mRNA"
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Homo sapiens interleukin IMAGE:7262176), complete BC069487 BC069487.1 GI:47480770
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/db_xref="HM:605619"
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GFSEIRGSVQAKDGNIDIRILRRTESLQDTKPANRCCLLRHLLRLYLDRVFKNYQTPD
HYTLRKISSLANSFLTIKKDLRLCHAHMTCHCGEEAMKKYSQILSHFEKLEPQAAVVK
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/clone_lib="NIH_MGC_244"
/note="Vector: PPCR-Script Amp_SK
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db_xref="GeneID:50604"
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1 (bases 1 to 531)

Strausberg RI, Feingold EA, Grouse LH, Derge JG, Klausner RD, Strausberg RI, Feingold EA, Grouse LH, Derge JG, Klausner RD, Strausberg RI, Feingold EA, Grouse LH, Derge JG, Klausner RD, Strausberg RI, Feingold EA, Schuler GD, Altschul SF, Zeeberg B, Buetow KH, Schaefer CF, Bhat NK, Hopkins RF, Jordan H, Moore T, Max SI, Wang J, Hsieh F, Diatchenko L, Marusina K, Farmer AA, Rubin GM, Hong L, Stapleton M, Soares MB, Bonaldo MF, Casavant TL, Scheetz TE, Brownstein MJ, Usdin TB, Toshiyuki S, Carninci P, Prange C, Raha SS, Loquellano NA, Peters GJ, Abramson RD, Mullahy SJ, Bosak SA, McEwan PJ, McKernan KJ, Malek JA, Gunaratne PH, SJ, Bosak SA, McEwan PJ, McKernan KJ, Malek JA, Gunaratne PH, SJ, Bosak SA, McEwan PJ, McKernan KJ, Malek JA, Gunaratne PH, SJ, Bosak SS, Worley KC, Hale S, Garcia AM, Gay LJ, Hulyk SW, Villalon E, Ketteman M, Madan A, Rodrigues S, Sanchez A, Whiting M, Helton E, Ketteman M, Madan A, Rodrigues S, Sanchez A, Whiting M, Madan A, Young AC, Shevchenko Y, Bouffard GG, Blakesley RW, Touchman JW, Green ED, Dickson MC, Rodriguez AC, Grimwood J, Schmutz J, Myers RM, Butterfield YS, Krzywinski MI, Skalaka U, Smailus DE, Schmerch A, Schein JE, Jones SJ and Marra MA.

Mammallan Gene Collection Program Team
Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Web site: http://www.hgsc.bcm.tmc.edu/cdna/
Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Loulseged,
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny, D.M., Nanavati,
A.N., Gibbs, R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tissue Procurement: Baylor Human Genome Sequencing Center cDNA Library Preparation: Baylor Human Genome Sequencing CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Baylor College of Medicine Human Genome
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Submitted (29-APR-2004) National Institutes of Health, Mammalian
Gene Collection (MGC), Cancer Genomics Office, National Cancer
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Email: cgapbs-r@mail.nih.gov
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the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
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                                                                                                                                                                                                                                                                                                                                                                                     PPCR-Script Amp SK(+)"
                                                                                                                                                                                                                                                                               IL10D, IL-20, MGC96907"
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RESULT 14 BC069523 Ś Ś Ş 밁 Ś 밁 Ś Ś 밁 밁 밁 밁 Ś 밁 Ś REFERENCE SOURCE KEYWORDS DEFINITION 맑 US-10-789-129-26 (1-144) x BC069487 Query Match: Percent Similarity: Best Local Similarity: Alignment Scores: VERSION ACCESSION Poccus AUTHORS ORGANISM 517 141 GluGluThrGlu 144 457 121 GluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet 397 101 HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu 337 277 217 157 81 AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys 100 61 41 97 21 Hominidae; Homo. 1 (bases 1 to 531) 1 (bases 1 to 531) Strausberg RL, Feingold EA, Grouse LH, Derge JG, Klausner RD, Strausberg RL, Feingold EA, Grouse LH, Derge JG, Klausner RD, Strausberg RL, Feingold EA, Grouse LH, Derge JG, Altschul SF, Zeeberg Collins FS, Wagner L, Shenmen CM, Schuler GD, Altschul SF, Zeeberg CD, Hiseher CF, Bhat NK, Hopkins RF, Jordan H, Moore T, Max SI, Wang J, Hsieh F, Diatchenko L, Marusina K, Farmer AA, Rubi CM, Hong L, Stapleton M, Soares MB, Bonaldo MF, Casavant TL, Scheetz TE, Brownstein MJ, Usdin TB, Toshiyuki S, Carninci P, Prange C, Raha SS, Loquellano NA, Peters GJ, Abramson RD, Mullahy SJ, Bosak SA, McEwan PJ, McKernan KJ, Malek JA, Gunaratne PH, Richards S, Worley KC, Hale S, Garcia AM, Gay LJ, Hulyk SW, Villalon DK, Muzny DM, Sodergren EJ, Lu X, Gibbs RA, Fahey J, Eukaryota; Metazoa; Mammalia; Eutheria; Homo sapiens Homo sapiens int IMAGE:7262093), Homo sapiens (human) BC069523 BC069523.1 GI:47479431 ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln GAGGAGACAGAA 528 GAACCTCAGGCAGCAGTTGTGAAGGCTTTGGGGGGAACTAGACATTCTTCTGCAATGGATG GCCAATTCCTTTACCATCAAGAAGGACCTCCGGCTCTGTCATGCCCACATGACATGC AGGGTATTTAAAAACTACCAGACCCCTGACCATTATACTCTCCGGAAGATCAGCAGCCTC ArgValPheLysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIleSerSerLeu GACACAAAGCCTGCGAATCGATGCTGCCTCCTGCGCCATTTGCTAAGACTCTATCTGGAC AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp GTGCAAGCCAAAGATGGAAACATTGACATCAGAATCTTAAGGAGGACTGAGTCTTTGCAA TGTGTGATCGCCACAAACCTTCAGGAAATACGAAATGGATTTTCTGAGATACGGGGCAGT 156 CysValIleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer sapiens interleukin E:7262093), complete 100.00% 99.31% 99.47% 4.73e-72 749.00 Chordata; Craniata; Vertebrata; Euarchontoglires; Primates; Cata (1-531)531 bp 20, mRNA cds. Conservative: Mismatches: Indels: Gaps: mRNA linear PRI (cDNA clone MGC:96884 531 143 0 ata; Euteleostomi; Catarrhini; PRI 08-MAR-2005 Zeeberg

456 120 336 80 276 60 216

40

REMARK

PUBMED REFERENCE AUTHORS

JOURNAL

JOURNAL

CONSRTM TITLE

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Percent Similarity:
Best Local Similarity:
                               US-10-789-129-26 (1-144)
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Proc. Natl. Acad. Sci. U.S.A. 99 (26), 16899-16903 (2007)
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Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Loulseged,
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny, D.M., Nanavati,
A.N., Gibbs, R.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequencing Center
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Contact: MGC help desk
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12. 1 (bases 1 to 531)

Strausberg RL, Feingold EA, Grouse LH, Derge JG, Klausner RD, Collins FS, Wagner L, Shenmen CM, Schuler GD, Altschul SF, Zeeberg B, Buetow KH, Schaefer CF, Bhat NK, Hopkins RF, Jordan H, Moore T, Max SI, Wang J, Hsieh F, Diatchenko L, Marusina K, Farmer AA, Rubin GM, Hong L, Stapleton M, Soares MB, Bonaldo MF, Casavant TL, Scheetz TE, Brownstein MJ, Usdin TB, Toshiyuki S, Carninci P, Prange C, Raha SS, Loquellano NA, Peters GJ, Abramson RD, Mullahy SJ, Bosak SA, McEwan PJ, McKernan KJ, Malek JA, Gunaratne PH, SJ, Bosak SA, McEwan PJ, McKernan KJ, Malek JA, Gunaratne PH, SJ, Bosak SA, Wczyn DM, Sodergren EJ, Lu X, Gibbs RA, Fahey J, Helton E, Ketteman M, Madan A, Rodrigues S, Sanchez A, Whiting M, Villalon DK, Muzny DM, Sodergren EJ, Lu X, Gibbs RA, Fahey J, Helton E, Ketteman M, Madan A, Rodrigues S, Sanchez A, Whiting M, Madan A, Young AC, Shevchenko Y, Bouffard GG, Blakesley RW, Touchman JW, Green ED, Dickson MC, Rodriguez AC, Grimwood J, Schmutz J, Myers RM, Butterfield YS, Krzywinski MI, Skalska U, Smailus DE, Schnerch A, Schein JE, Jones SJ and Marra MA.

11. Schmutz J, Myers RM, Butterfield YS, Krzywinski MI, Skalska U, Smailus DE, Schnerch A, Schein JE, Jones SJ and Marra MA.

12. Droc Marl Acid Sci II S A Go [26] 1680a-16903 (2002)
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Direct Submission

Direct Submission

Submitted (29-APR-2004) National Institutes of Health, Mammalian Submitted (29-APR-2004) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Gene Collection (MGC), Cancer Genomics Office, National Cancer Gene Collection (MGC), Cancer Genomics Office, National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National Cancer General National National Cancer General National Cancer General National Nati
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                                                                                                                                                          Director MGC Project.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BC069559.1 GI:47480991
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FEATURES

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Best Local Similarity:
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Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Loulseged,
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny, D.M., Nanavati,
A.N., Gibbs, R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Baylor Human Genome Sequencing Center
cDNA Library Preparation: Baylor Human Genome Sequencing Center
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Baylor College of Medicine Human Genome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Clone distribution: MGC clone distribution information can be for through the I.M.A.G.B. Consortium/LLNL at: http://image.llnl.gov Series: IRBR Plate: 1 Row: b Column: 3.

Location/Qualifiers
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Center code: BCM-HGSC
                                                                                                                                                                            AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp 60
                                                                                                                                                                                                                                                          ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln 40
                                                                                                                                                                                                                                                                                                                            CysValIleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer
                      AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys 100
                                                                AGGGTATTTAAAAACTACCAGACCCCTGACCATTATACTCTCCGGAAGATCAGCAGCCTC
                                                                                     ArgValPheLysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIleSerSerLeu 80
                                                                                                                                                 GACACAAAGCCTGCGAATCGATGCTGCCTCCTGCGCCCATTTGCTAAGACTCTATCTGGAC
                                                                                                                                                                                                                            GTGCAAGCCAAAGATGGAAACATTGACATCAGAATCTTAAGGAGGACTGAGTCTTTGCAA
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/tissue_type="PCR rescued clones"
/clone_Tib="NIH_MGC_244"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ALGELDILLQWMEETE"
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/db_xref="MIM:605619"
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Conservative:
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           141 GluGluThrGlu 144
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                                             GluProGlnAlaAlaValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet
                                                                  HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu 120
GAGGAGACAGAA
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Search completed: December 28, 2005, 15:08:47 Job time : 2947 secs

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Result
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-UNITS=bite -STRATT=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
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-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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-MODEL=frame+_p2n.model
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1: geneseqn198
2: geneseqn199
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7: geneseqn200
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Adm95042 Human Int
Adv42800 Human psy
Aax77692 Human Zcy
Acf05161 Polynucle
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ACD25355 ACF00404 ACA60431	ACF13	ACF19799 ACD22087		ACD81609			ACA 1368		ADY3195			ABS74446			AAS46229		AAS21292		AAA						12 ADL15289		AEB2				3 ADS18362			2 ADQ8806	0 ADG4666	0 AAD6
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ALIGNMENTS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New microarray comprising probes for genes involved in psychoneuroendocrinimmune (PNI) activity, useful in diagnosing condition associated with PNI activity, e.g., inflammatory or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       microarray; psychoneuroendocrinimmune; chronic fatigue; non-insulin dependent diabetes; allergy; immune disorder; cancer; neoplasm; infection; expressed sequence tag; ss.
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                   CC This invention describes a novel mammalian four alpha helix cytokine CC designated Zcytol0 which has vulnerary, cytostatic, anti-inflammatory, CC anti-asthmatic and haemostatic activity. Zcytol0 may be a growth and/or CC maintenance factor in the trachea and salivary glands, stomach, pancreas CC and muscle. Zcytol0 polynucleotides are useful as sources of primers and CC probes, and to determine if the Zcytol0 gene is present on chromosome 1, CC or to detect any mutations that may have occurred. Zcytol0 cc polynucleotides are also useful as gene therapy reagents. The Zcytol0 cc polypeptide, its fragments, and antibodies, as well as compounds cc identified as having binding affinity to Zcytol0, should be useful in the treatment of conditions associated with abnormal physiology or CC creatment of conditions associated with abnormal physiology or CC or degenerative conditions or altered immunity. The products are also CC useful for modulating cell proliferation, cell differentiation or CC cytokine production in the prevention or treatment of conditions or altered immunity. The products are also CC cytokine production in the prevention or treatment of conditions are cytokine production. The proliferation, cell differentiation or cytokine production.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Inhibiting the growth and/or proliferation of cervical cancer cells, useful for treating cervical cancer or human papilloma virus infection, comprises bringing Interleukin-20 (IL-20) into contact with the cervical cancer cells.
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AAD61821
Human cytokine-like polypeptide-10 (Zcyto10) cDNA
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autoimmune disease; insulin-dependent diabetes mellitus; immune response; rheumatoid arthritis; multiple sclerosis; infection; radiation therapy; asthma; bronchitis; wound healing; periodontal disease; antiinflammatory; psoriasis; eczema; dry skin; oral cavity; muscle atrophy; haemostatic; cancer; vulnerary; cytokine-like polypeptide-10; Zcyto10; therapy; thrombocytopaenia; mune disease; insulin-dependent diabetes mellitus; immune response; chromosome 1; gene; ss.

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Novel human cytokine-like polypeptide-10, Zcyto10, useful for treating insulin-dependent diabetes mellitus, rheumatoid arthritis, multiple sclerosis, asthma, psoriasis and inhibiting cancer cell growth.

Disclosure; Col 29-30; Opp; English

The invention relates to an isolated human cytokine-like polypeptide-10 (Zcyto10) polypeptide. Zcyto10 is useful for prevention or treatment of CC conditions characterised by improper cell proliferation, cell CC differentiation or cytokine production. Zcyto10 is useful for treating CC autoimmune diseases such as insulin-dependent diabetes mellitus, CC rheumatoid arthritis, multiple sclerosis, etc., by inhibiting cellular CC immune response. The invention is useful for inhibiting cellular CC growth or proliferation, for stimulating immune system to combat CC microbial or viral infections, for increasing platelet production, and CC thus useful in cancer patients who experience thrombocytopaenia due to CC cancer therapy or radiation therapy. Zcyto10 is useful in tracheacy specific or tracheobronchial-specific applications, such as in the CC underlying the same, in regulating mucous production or mucocillary CC clearance of debris or in treatment of asthma, bronchits or other CC diseases of the tracheobronchial tract. Zcyto10 may also enhance wound the same in the treatment of affected tissues which may be useful to the treatment of periodontal disease. Zcyto10 can be useful to treat skin conditions such as psoriasis, eczema and dry skin in CC used to treat skin conditions such as psoriasis, eczema and dry skin in þe

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                                                                                                              cytokine-like polypeptide-10; Zcyto10; wound healing; platelet proliferation; wound repair; mucous production; asthma; bronchitis; tracheobronchial tract disease; periodontal disease; skin condition; psoriasis; eczema; dry skin; protein therapy; human; vulnerary; respiratory; antibronchitic; dental; dermatological; gene;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to novel mammalian cytokine-like polypeptide-10 (Zcytol0) proteins and polynucleotides encoding such proteins. Sequences of the invention are useful for promoting the healing proteins. Sequences of the invention are useful for promoting the healing of wounds and for stimulating the proliferation of platelets. In of wounds and for stimulating the proliferation of platelets. In particular Zcytol0 polypeptides may be used in trachea-specific or tracheobronchial-specific applications such as maintenance or wound repair of the tracheobronchial epithelium or cells underlying it; in regulating mucous production; mucocilary clearance of debris; in tracheobronchial tract. They are also used to promote regeneration of affected tissues which may be especially useful in the treatment of affected tissues which may be especially useful in the treatment of periodontal disease. Zcytol0 polypeptides can be used to treat skin conditions such as psoriasis, eczema and dry skin in general. They are also used in protein therapy. The present sequence is human Zcytol0 DNA.
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25-NOV-1998;
17-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated mammalian cytokine-like polypeptide-10 (Zcytol0), ufor treating asthma, bronchitis and other tracheobronchial damage
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GCCAATTCCTTTCTTACCATCAAGAAGGACCTCCGGCTCTGTCATGCCCACATGACATGC
              AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
                                                    AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp
                                                                                                                                                                GTGCAAGCCAAAGATGGAAACATTGACATCAGAATCTTAAGGAGGACTGAGTCTTTGCAA
                                                                                                                                                                                ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln
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26-NOV-1997;
25-NOV-1998;
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                             New ZcytolO polypeptides and polynucleotides, useful for treating autoimmune diseases (e.g. insulin dependent diabetes mellitus, multiple sclerosis or rheumatoid arthritis), or skin conditions (e.g. psoriasis)
                                                                                                                                                                                                                                                                                                                                                                                                      07-OCT-2004
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         Disclosure;
                                                                    WPI; 2004-552564/53.
                                                                                 Conklin DC,
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98US-00199586.
2003US-00413661.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention provides mammalian cytokine-like polypeptide-10 (2cyto10) proteins and their encoding polynucleotides. The invention is useful for treating conditions characterised by improper cell proliferation, cell differentiation or cytokine production, specifically for treating autoimmune diseases such as insulin dependent diabetes mellitus, multiple sclerosis and rheumatoid arthritis and for inhibiting cancer cell growth. The invention is also used to stimulate the immune cystem, to enhance wound healing and promote regemeration of affected tissues, to increase platelet count and to treat asthma and skin conditions such as psoriasis, eczema and dry skin. The invention acts as an antidiabetic, neuroprotective, antirheumatic, antiarthritic, cytostatic, immunostimulant, vulnerary, antiasthmatic, antipsoriatic and cermatological agent. The invention is also used in gene therapy. The present sequence is human Zcytol0 longer form DNA.
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standard; DNA; 926
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Cytokine-like polypeptide-10; Zcyto10; therapy; asthma; infection; psoriasis; eczema; dry skin; wound healing; platelet proliferation;

Human 21-OCT-2004

cytokine-like polypeptide-10 (Zcyto10) long

form DNA (first

entry)

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25-NOV-1998;
15-APR-2003;
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                                                                                                                                                                                                                                                                                                       Sequence 926
                                                                                                                                                                                                                                                                                                                                                                                                       New isolated antibody that binds to a ZcytolO polypeptide, useful treating asthma, microbial or viral infections, and for promoting healing of wounds, psoriasis, eczema or dry skin.
                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 1; 32pp; English.
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         AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
                                                  ArgValPheLysAsnTyrGlnThrProAspHisTy
                                                                                           AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp
                                                                                                                   GTGCAAGCCAAAGATGGAAACATTGACATCAGAATCTTAAGGAGGACTGAGTCTTTGCAA
                                                                                                                                                            TGTGTGATCGCCACAAACCTTCAGGAAATACGAAATGGATTTTCTGACATACGGGGCAGT
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98US-00199586.
2003US-00413661.
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Percent Similarity:
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Query Match:
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ADR24325
US-10-789-129-26
                                                                                                                                     Alignment
                                                                                                                                                                                                     The invention relates to a method of classifying a breast cancer patient according to prognosis by determining the similarity between the level of expression of each of five genes for which markers are listed in the specification, in a cell sample taken from the breast cancer patient, to control levels of expression for each respective five genes to obtain a patient similarity value. The methods are useful for classifying a breast cancer patient according to prognosis. Kits and computer program products are useful for data analysis using the diagnostic, prognostic and statistical methods of the invention. This sequence corresponds to a marker used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                               Classifying a breast cancer patient according to prognosis comprises determining the similarity between the level of expression of each of five genes in a cell sample taken from patient, to control levels.
                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 186; 226pp; English.
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RESULT 11
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AEA50048 sta AEA50048; 11-AUG-2005 Full length

IL-20 coding sequence

(first

entry)

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Percent Similarity:
Best Local Similarity:
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P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antibody that selectively binds to a Zcytol0 polypeptide useful modulating cell proliferation, cell differentiation or cytokine production in the prevention or treatment of e.g. autoimmune disease
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This sequence encodes full length interleukin-20 (IL-20), longer form IL-CC 20x1. The method of the invention for producing an antibody to IL-20 comprises inoculating an animal with an IL-20 polypeptide which elicits CC an immune response in the animal to produce the antibody, and isolating CC the antibody from the animal. The antibody is useful for treating a CC pathological condition in a subject associated with IL-20 activity such CC disease, ulcerative colitis, Crohns' disease, arthritis, atopic CC dermatitis and psoriasis, or acute inflammatory condition such as CC endotoxemia, septicemia, toxic shock syndrome and infectious disease. An CC inflammatory disease in which IL-20 plays a role such that the CC inflammatory disease in which IL-20 plays a role such that the CC inflammatory disease in which IL-20 plays a role such that the CC inflammatory is reduced, where the antagonist comprises an antibody, CC antibody fragment, or binding polypeptide that specifically binds a CC polypeptide or polypeptide fragment of IL-20, and where the inflammatory activity of IL-CC 20 is reduced. The method is useful for producing such therapeutic CC antibody. The antibody or antibody fragment, produced using the above CC method, binds to IL-20, and inhibits or reduces the pro-inflammatory activity of IL-20. IL-20-induced proliferation or differentiation of inhibited minimater has antibody antibody fragment of IL-20 and antibody fragment in antibody in the antibody or antibody fragment antibody such therapeutic CC method, binds to IL-20 and inhibits or reduces the pro-inflammatory inhibited minimater has antibody antibody fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Producing an antibody to an interleukin-20 (IL-20) polypeptide, useful for treating inflammatory diseases e.g., Crohn's disease, by inoculating an animal with the polypeptide eliciting an immune response to produce
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Percent Similarity:
Best Local Similarity:
Query Match:

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel method for producing an antibody to a coplypeptide. The method comprises inoculating an animal with a coplypeptide selected from any of SEQ ID NO: 14, 15, 17 or 19 as given in the specification, where the polypeptide elicits an immune response in the animal to produce the antibody, and isolating the antibody from the canimal. The antibody specifically binds to the pro-inflammatory cytokine interleukin 20 (IL-20 or Zcytol0), or one its receptors IL-20RA (ZcytpR7) cor II-20RB (pDIRS1). The method of the invention demonstrates and antiminflammatory, gastrointestinal-gen., antiplicary antiarthritic, cantimicrobial applications. The antibody may be useful for treating a cromatological condition in a subject associated with IL-20RA, such as a chronic inflammatory condition selected from inflammatory bowel disease, chronic inflammatory condition selected from inflammatory bowel disease, culcerative colitis, Crohn's disease, arthritis, aropic dermatitis and conditions, concer, or an acute inflammatory condition such as collected sequence is that of a human IL-20 precursor cDNA clone of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Producing an antibody to an interleukin 20RA (IL-20RA) polypeptide, useful for treating inflammatory diseases e.g., arthritis, by inoculating an animal with the polypeptide which elicits an immune response to produce the antibody.
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                                                                                                                                                                                                                                                                                                                                                                                                             The invention provides a new purified polynucleotide that comprises the concleic acid sequence A-B-C-D-E; where: A- start codon, B- polynucleotide concleic acid sequence of 13 nucleic acids, where the sequence has homology with 13 concleic pair (bp) sequence, where at least 5 nucleic acids are identical to code for a stop codon, C- polynucleotide sequence comprising 10, 13, 16 concleic acids of a 19 bp sequence (ABY75139), and where the nucleic acid sequence does not code for a stop codon, C- polynucleotide sequence comprising 10, 13, 16 concleic acids of a 19 bp sequence (ABY75140), D- polynucleotide concleic acids of a 19 bp sequence (ABY75140), D- polynucleotide concleic acid from 9 sequences (ABY75144-150) where at least 50% nucleotides are either adenine or thymine, and where only E encodes a stop codon. The colynucleotide is useful for producing a nucleic acid construct for concression of an amino acid sequence of interest. The present sequence corpresents a human interleukin-20 (IL-20) codon optimised fragment for construct for expression in E. coli
                                                                                                                                                                                                                              US-10-789-129-26 (1-144) x ABV75151 (1-462)
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                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 36; 40pp; English
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ArgValPheLysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIleSerSerLeu
                                                                 AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp
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                                                                                                                                                            The invention relates to Sym 1 and Sym 2 tight regulated synthetic promoters and methods for enhancing the expression of recombinant proteins. The Sym 1 and Sym 2 tight regulated synthetic promoters are useful for controlling uninduced, leaky expression of proteins that are toxic and interfere with production of recombinant protein from an Escherichia coli expression system. They are useful for controlling basal expression of target genes. The present sequence is the human interleukin [II]-20 DNA. This sequence is used in the comparison of promoter
                                                                                               Sequence 462
                                                                                                                                                                                                                                                                                                                                                                                                                                                      New Syn 1 and Syn 2 tight regulated synthetic promoters, useful for controlling uninduced, leaky expression of proteins that are toxic and interfere with production of recombinant protein from an Escherichia coli
                                                                                                                                                                                                                                                                                                                                                                                         Example 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                         expression
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DB; AEB26650.
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                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                       system.
                                                                                             BP; 121 A; 131
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                     New polynucleotide, useful for producing a nucleic acid expression of an amino acid sequence of interest.
Example 1; Page 37; 40pp; English.
                                                                           Pownder TA,
                                                                                                                       13-APR-2001; 2001US-0283688P
                                                                                                                                              12-APR-2002; 2002WO-US011513
                                                                                                                                                                     24-OCT-2002
                                                                                                                                                                                         WO200283853-A2
                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                           Protein synthesis; data mining; recombinant; interleukin-20; IL-20;
                                                                                                                                                                                                                                                                                EC sequence
                                                                                                                                                                                                                                                                                                        19-FEB-2003
                                                                                                                                                                                                                                                                                                                               ABV75154;
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                                                                                                  ( OMYZ)
                                                      2003-067576/06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CysValIleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer
                                                                                                                                                                                                                                                                                                                                                                                                  GAGGAGACCGAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention provides a new purified polynucleotide that comprises the concleic acid sequence A-B-C-D-E; where: A= start codon, B= polynucleotide code sequence of 13 nucleic acids, where the sequence has homology with 13 code base pair (bp) sequence, where at least 5 nucleic acids are identical to code for a stop codon, C= polynucleotide sequence comprising 10, 13, 16 code for a stop codon, C= polynucleotide sequence comprising 10, 13, 16 code for a stop codon, C= polynucleotide sequence comprising 8 bp (ABV75141), B= polynucleotide sequence selected from 9 sequences (ABV75141), B= polynucleotide sequence selected codenine or thymine, and where only E encodes a stop codon. The codonucleotide is useful for producing a nucleic acid construct for coplynucleotide is useful for producing a nucleic acid construct for coplynucleotide sequence comprising an expression cassette (EC) polynucleotide sequence comprising an expression cassette (EC) colynucleotide sequence inserted upstraman interleukin-20 (IL-Ca) coden continued for producing a nucleic acid construct for colons continued for producing an expression cassette (EC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 513 BP; 143 A; 145 C;
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                                                                                                                                                                                                HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu 120
                                                                                                                                                                                                                                                                                                                                 AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys 100
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                                            GluGluThrGlu 144
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-MODBL-frame+ p21.model -DEV=xlp
-Q-/cgn2 1/USPTO_spool_p/US10789129/runat_23122005_152720_20276/app_query.fasta_1.327
-Q=/cgn2 1/USPTO_spool_p/US10789129/runat_23122005_152720_20276/app_query.fasta_1.327
-DB=EST -QFWT=fastap -SUFFIX=p21.rst -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS=bits -START=1 -ENDE-1 -MATRIX=blosum62 -TRAUS=humah40.cdi -LIST=45
-DOCALIGN=200 -THR SCORE=pct -THR MAX=100 -THR MIN=0 -ALIGN=15 -MODE=LOCAL
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-USER=US10789129 @GCN 1 _8010 @Funnat_23122005 _152720 _20276 -NCPU=6 -ICPU=3
-NO MMAP -LARGEQUERY -NEG SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
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ALIGNMENTS

ĀL		JOURNAL PUBMED REFERENCE AUTHORS	TITLE	REFERENCE AUTHORS	ACCESSION VERSION KEYWORDS SOURCE ORGANISM	RESULT 1 AY402693 LOCUS DEFINITION
Submitted (16.NOV-2003) Celera Genomics, 45 West Gude Drive, Submitted (16.NOV-2085), USA Rockville, MD 20850, USA This sequence was made by sequencing genomic exons and ordering	Todd,M.A., Tanenbaum,D.M., Civello,D.R., Lu,F., Murphy,B., Ferriera,S., Wang,G., Zheng,X.H., White,T.J., Sninsky,J.J., Adams,M.D. and Cargill,M. Direct Submission	gene trios Science 302 (5652), 1960-1963 (2003) 14671302 2 (bases 1 to 531) Clark,A.G., Glanowski,S., Nielson,R., Thomas,P., Kejariwal,A.,	Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M. Inferring nonneutral evolution from human-chimp-mouse orthologous	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 531) Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A.,	ĀY402693 AY402693.1 GI:39758676 GSS. Homo sapiens (human) Homo sapiens	AY402693 531 bp DNA linear GSS 15-DEC-2003 Homo sapiens IL20 gene, VIRTUAL TRANSCRIPT, partial sequence, genomic survey sequence.

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                       Hominidae; Pan.

1 (Dases 1 to 531)

1 (Dases 1 to 531)

Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
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Clark,A.G., Glanowski,S., Nielson,R., Thomas,P., Kejariwal,A., Todd,M.A., Tanenbaum,D.M., Civello,D.R., Lu,F., Murphy,B., Ferriera,S., Wang,G., Zheng,X.H., White,T.J., Sninsky,J.J., Adams,M.D. and Cargill,M.
Direct Submission
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Mus musculus IL20 gene, VIRTUAL
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/gene="IL20"
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/mol_type="genomic DNA"
/db_xref="taxon:9598"
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Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
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               Direct Submission

Submitted (16-APR-2002) Yoshinide Hayashizaki, The Institute of Physical and Chemical Research (RIKEN), Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center (GSC), RIKEN Yokohama Institute; 1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama RIKEN Yokohama Institute; 1-7-22 Suehiro-cho, Tsurumi-ku, Yokoham Kanagawa 230-0045, Japan (E-mail:genome-res@gsc.riken.jp, URL:http://genome.gsc.riken.jp/, Tel:81-45-503-9222, Pax:81-45-503-9216)
                                                                                                                                                                                      Adachi, J., Aizawa, K., Akimura, T., Arakawa, T., Bono, H., Carninci, P. Pukuda, S., Furuno, M., Hanagaki, T., Hara, A., Hashizume, W., Hayashidak, K., Hayatsu, N., Hiramoto, K., Hiramoka, T., Hirozane, T., Harashidak, K., Hayatsu, N., Hiramoto, K., Hiramoka, T., Hirozane, T., Hori, F., Imotani, K., Ishii, Y., Itoh, M., Kagawa, I., Kasukawa, T., Katoh, H., Kawai, J., Kojima, Y., Kondo, S., Konno, H., Kouda, M., Koya, S., Kurihara, C., Matsuyama, T., Miyazaki, A., Murata, M., Nakamura, M., Nishi, K., Nomura, K., Numazaki, R., Ohno, M., Ohsato, N., Okazaki, Y., Saito, R., Saitoh, H., Sakai, C., Sakai, K., Sakazume, N., Sano, H., Sasaki, D., Shibata, K., Shinagawa, A., Shiraki, T., Sogabe, Y., Tagama, M., Tagawa, A., Takahashi, F., Takaku-Akahira, S., Takada, Y., Tanaka, T., Tomaru, A., Toya, T., Yasunishi, A., Paranasani, M., and Hayashizaki, Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      The FANTOM Consortium and the RIKEN Group Phase I & II Team.
Analysis of the mouse transcriptome of 60,770 full-length cDNAs
Nature 420, 563-573 (2002)
6 (bases I to 1189)
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High-efficiency full-length cDNA cloning
Meth. Enzymol. 303, 19-44 (1999)
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Please visit our web site for further details.
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/clone_Iib="RIKEN full-length enriched mouse cDNA library"
/dev_stage="adult"
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/db_xref="FANTOM_DB:7530415H19"
/db_xref="taxon:10090"
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/mol_type="mRNA"
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Contact: Yoshihide Hayashizaki
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Computer-based methods for the mouse full-length cDNA encyclopedia: real-time sequence clustering for construction of a nonredundant cDNA library. Genome Res. 11 (2), 281-289 (2001)

Kondo, S., Shinagawa, A., Saito, T., Kiyosawa, H., Yamanaka, I., Aizawa, K., Fukuda, S., Hara, A., Itoh, M., Kawai, J., Shibata, K. and Hayashizaki, Y.
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The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-
Tel: 81-45-503-9222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Computational Analysis of Full-Length Mouse cDNAs Compared with Human Genome Sequences. Mamm. Genome. 12, 673-677 (2001)
Please visit our web site (http://genome.gsc.riken.go.jp) for
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RIKEN integrated sequence analysis (RISA) system--384-format sequencing pipeline with 384 multicapillary sequencer. Genome 10 (11), 1757-171 (2000)
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                                                                                                                                                                                                                                                                                                           /tissue_type="eyeball"
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/lab_host="DH10B"
/clone_lib="RIKEN full-length enriched, adult male
eyeball"
/note="Site 1: SalI; Site 2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to prepare mouse tissues. 1st strand cDNA was
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                                           RZPD Deutsches Ressourcenzentrum fuer Heubnerweg 6, D-14059 Berlin, Germany Email: www.rzpd.de
                                                                                   Human T-Lymphocytes library
Unpublished (2005)
Contact: Inge Arlart
RZPDLIB; (Human T-Lymphocytes) RZPD LIB No.9016
http://www.rzpd.de/cgi-bin/products/set.cgi?libNo=9016 Contact:
                                                                                                                                                                          Hominidae; Homo
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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RZPD Deutsches Ressourcenzentrum fuer Genomforschung
Heubnerweg 6, D-14059 Berlin, Germany
Tel: +49 30 32639 100
Fax: +49 30 32639 111
 GluGlu 142
                                                           GluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet 140
                                                                                                                  HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
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                              GAGGTCCACGCTGCCATTAAATCCCTGGGAGAGCTCGACGTCTTTCTAGCCTGGATT
                                                                                        CÁCTGCAGGCAGGAÁGCCACCAATGCCACCAGAGTCÁTCCATGACAACTATGATCAGCTG
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/clone="RZPDD9016M1629"
/tissue_type="T-Lymphocytes"
/dev_stage="adult"
/lab_host="PH10B"
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/db_xref="taxon:9606"
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Hara,A., Hiramoto,K., Hori,F., Ishii,Y., Ito,M., Kawai,J.,
Konno,H., Kouda,M., Koya,S., Mateuyama,T., Miyazaki,A., Nomura,K.,
Ohno,M., Okazaki,Y., Okido,T., Saito,R., Sakai,C., Sakai,K.,
Sano,H., Sasaki,D., Shibata,K., Shinagawa,A., Shiraki,T.,
Sogabe,Y., Suzuki,H., Tagami,M., Tagawa,A., Takahashi,F.,
Takeda,Y., Tanaka,T., Toya,T., Muramateu,M. and Hayashizaki,Y.
RIKEN Mouse ESTs (Arakawa,T., et al. 2001)
Unpublished (2001)
On Jun 14, 2000 this sequence version replaced gi:8522215.
Contact: Yoshihide Hayashizaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: genome-res@gsc.riken.jp, URL:http://genome.gsc.riken.jp/
Carninci, P., Shibata, Y., Hayatsu, N., Sugahara, Y., Shibata, K.,
Itoh, M., Konno, H., Okazaki, Y., Muramatsu, M. and Hayashizaki, Y.
Normalization and subtraction of cap-trapper-selected cDNAs to
prepare full-length cDNA libraries for rapid discovery of new
genes. Genome Res. . 10 (10), 1617-1630 (2000)
wagi, K., Pujiwake, S., Inoue, K., Togawa, Y., Izawa, M., Ohara, B.,
Watahiki, M., Yoneda, Y., Ishikawa, T., Ozawa, K., Tanaka, T.,
Matsuura, S., Kawai, J., Okazaki, Y., Muramatsu, M., Inoue, Y., Kira, A.
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Computer-based methods for the mouse full-length cDNA encyclopedia: real-time sequence clustering for construction of a nonredundant cDNA library. Genome Res. 11 (2), 281-289 (2001) Kondo, S., Shinagawa, A., Saito, T., Kiyosawa, H., Yamanaka, I., Aizawa, K., Fukuda, S., Hara, A., Itoh, M., Kawai, J., Shibata, K. and Hayashizaki, Y.
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The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi; Muroidea; Muridae; Murinae; Mus.
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Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human Genome Sequences Mamm. Genome. 12, 673-677 (2001)

Please visit our web site (http://genome.gsc.riken.go.jp/) for further details.
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                                    /tissue_type="eyeball"
/dev_stage="adult"
/lab_host="DH10B"
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'clone_lib="RIKEN full-length enriched, adult male
                                                                                                                                        sex="male"
                                                                                                                                                                                     clone="7530415H19"
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/note="Site 1: SalI; Site 2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to prepare mouse tissues. 1st strand cDNA was
                                                                                                                                                                                                  primed with a primer [5'
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Pred. Alignment

Scores:

Ś 밁 Ś 밁 S 밁 á 밁 S 밁 RESULT US-10-789-129-26 (1-144) x BB064801 (1-695) Query Match: Best Local Similarity: Percent Similarity: 242 AGATGGATGGAGGAG 256 138 GlnTrpMetGluGlu 142 182 ATAGAGTTGGAACTTCAGGCÁGCGGTGGTAAAGGCTTTGGGÁAÁACTAGGCÁTTCTTCTG 118 GluLysLeuGluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeu 123 ATGGCATGTCA-TGTGGGGAAGAAGCAATGGAGAAATACAACCAAATTCTGAGTCACTTC 98 MetThrCysHisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPhe 63 78 SerSerLeuAlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHis 58 TyrLeuAspArgValPheLysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIle w AGCAGCCTCGCCAAATCCTTTCTTATCATCAAGAAGGACCTCTCAGTCTGTCATTCTCAC TATCTGGACAGGGTATTCAAAGTCTACCAGACCCCTGACCNCCATACCCTGAGAAAGATC 6.17e-30 328.00 38.24% 80.00% 43.56% Gaps: Mismatches: Indels: Conservative: 181 117 122 97 62

JOURNAL COMMENT

Unpublished (2005) Contact: Inge Arlart

RZPD Deutsches Ressourcenzentrum fuer Genomforschung GmbH Heubnerweg 6, D-14059 Berlin, Germany Email: www.rzpd.de

TITLE AUTHORS

and Korn, B.

(bases 1 to 699) T-Lymphocytes library

Schneider, D.

REFERENCE

CR992594 LOCUS

VERSION KEYWORDS

ACCESSION DEFINITION

mRNA sequence. CR992594

CR992594.1

GI:68286479

CR992594 RZPD no.9016 Homo sapiens cDNA clone

near EST 28-JUN-2005 RZPDp9016C1545 5',

SOURCE

ORGANISM

Homo sapiens

Homo sapiens (human)

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;

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Percent Similarity:
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Tel: +49 30 32639 100
Fax: +49 30 32639 111
www.rzpd.de
This clone is available
                                                                                                                                                                                                                                                                                                                                                                                                                                         21 ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln 40
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http://www.rzpd.de/cgi- bin/products/cl.cgi?CloneID=RZPDp9016C1545

contact RZPD (product- support@rzpd.de) for further information.

Primer name: qe3 4 , Primer sequence: CGGATAACAATTCACACAG.

Location/Qualifiers
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http://www.rzpd.de/cgi-bin/products/set.cgi?libNo-9016 Contact:
                                                                                                          HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
                                                                                                                                                                                                                                                                                                                      ATCATTAAGCCCTTAGATGTGTGCTGCGTGACCAAGAACCTCCTGGCGTTCTACGTGGAC
                       GluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMet 140
                                                                                                                                                                                              AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys 100
                                                                                                                                                                                                                                                                      ArgValPheLysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIleSerSerLeu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CysVallleAlaThrAsnLeuGlnGlulleArgAsnGlyPheSerAspIleArgGlySer
GAGGTCCACGCTGCCATTAAATCCCTGGGAGAGCTCGACGTCTTTCTAGCCTGGATT
                                                                           CACTGCAGGCAGGAAGCCACCAATGCCACCAGAGTCATCCATGACAACTATGATCAGCTG
                                                                                                                                                           GCCAACTCTTTCCTCTACATGCAGAAAACTCTGCGGCAATGTCAGGAACAGAGGCAGTGT
                                                                                                                                                                                                                                        AGGGTGTTCAAGGATCATCAGGAGCCCAAAACCCCCAAAATCTTGAGAAAAATCAGCAGCATT
                                                                                                                                                                                                                                                                                                                                                           AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp
                                                                                                                                                                                                                                                                                                                                                                                                        ATCCAAGCTAAGGACACCTTCCCAAATGTCACTATCCTGTCCACATTGGAGACTCTGCAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /dev_stage="adult"
/lab_host="DH10B"
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/tissue_type="T-Lymphocytes"
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/db_xref="taxon:9606"
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Tissue Procurement: Dr. Gary W. Hunninghake, U of I
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Clone distribution information can be obtained
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UI-H-FT1-bic-n-07-0-UI.81 NCI_CGAP_FT1 Homo sapiens cDNA clone
UI-H-FT1-bic-n-07-0-UI 3', mRNA sequence.
CA309757
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Seq primer: M13 FORWARD
POLYA=Yes.
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Unpublished (1997)
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1 (bases 1 to 762)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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                  the conditions, other donor macrophages in different conditions). The mRNA samples were pooled for library construction. Control 0 hours; control 3 hours; control 24 hours; LPS 100 ng/ml, 3 hours; LPS 100 ng/ml, 24 hours; PMA 10 ng/ml, 3 hours; MA 10 ng/ml, 24 hours; Klebsiella moi 10, 24 hours; Klebsiella moi 10, 24 hours; Klebsiella moi 10, 24 hours; Klebsiella vector (Ad5 CMV eGFP), moi 500, 3 hours; Adenoviral vector (Ad5 CMV eGFP), moi 500, 24 hours; Adenoviral vector (Ad5 CMV eGFP), moi 500, 24 hours; Adenoviral vector (Ad5 CMV eGFP), moi 500, 24 hours; Adenoviral vector that it adenovirus moi 500, 3 hours; Adenovirus moi 500, 3 hours; Adenovirus moi 500, 24 hours; The library was normalized according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806, 1996. First strand cDNA synthesis was primed with an oligo-dT primer containing a Not I site. Double stranded cDNA was ligated to an EcoR I adaptor, digested with Not I, and cloned directionally into pT773-Pac vector. The oligonucleotide used to prime the synthesis of first-strand cDNA contains a library tag sequence that is located between the Not I site and the (dT)18 tail. The sequence tag for this library is GGCCATGCCG. The tissue was provided by Dr. Gary W. Hunninghake of the University of Iowa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone lib="NCI_CGAP_FT1"
/note="Organ: Lung; Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site_1: EcoR I; Site_2: Not I; NCI_CGAP_FT1 is a normalized cDNA library constructed from a pool of 81 RNA samples from Alveolar Macrophages challenged with different treatments. The mRNA samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               were a mixture of these conditions (times refer to incubations following isolation by bronchoalveolar lavage) (some normal donor macrophages were cultured in some of
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/lab_host="DH10B (Life Technologies)"
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/clone="UI-H-FT1-bic-n-07-0-UI"
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   TISSUE=Human
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Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W.Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and Simpson,A.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BF738279.1 GI:12065059
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PM4-KT0045-151200-006-g02
BF738279
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?ti=PM4&t2=PM4-KT0045-
151200-006-g02&t3=2000-12-15&t4=1)
                                                                                                        Brazil
Tel: +55-11-2704922
                                                                                                                                                       Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                          Shotgun sequencing of the human transcriptome with ORF
                                                                                                                                          Rua Prof. Antonio Prudente 109, 4 andar, 01509-010,
                                                                                                                                                                                                                                          sequence tags
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TAG_SEQ=GGCCATGCCG"
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ens cDNA,
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source

/organism="Homo sapiens" /mol_type="mRNA" /db_xref="taxon:9606"

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REFERENCE
AUTHORS
TITLE
                                                                                                                                                                                                           JOURNAL COMMENT
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KEYWORDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CA306526 748 bp mRNA linear UI-H-FT1-bhw-m-10-0-UI.sl NCI_CGAP_FT1 Homo sapiens UI-H-FT1-bhw-m-10-0-UI 3', mRNA sequence.
                                                        Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. Gary W. Hunninghake, U of I
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Clone distribution information can be obtained
from Dr. M. Bento Soares, bento-soares@uiowa.edu
                                                                                                                                                                                                                                                                                                                                                        Bukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Euarchontoglires; Primates; Cata
                                                                                                                                                                                                                               Unpublished (1997)
                                                                                                                                                                                                                                                                                                                Hominidae; Homo.
1 (bases 1 to 748)
                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (human)
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                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                         NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                         Seq primer: M13 FORWARD
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                                                                                                                                                                                                                                                   Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Torgan: bladder tumor; Vector: puc18; Site_1: SmaI; Site_2: SmaI; A mini_Tibrary was made by cloning products derived from ORBSTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /dev_stage="Adult"
/clone_lib="KT0045"
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/mol_type="mRNA"
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Matches:
Conservative:
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Indels:
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                                                                                                                                                                                                                                                                             Anatomy
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                                                                                                                                                                                                                                                                             Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                  Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cDNA clone
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                                                                                       be obtained
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US-10-789-129-26 (1-144) x CA306526 (1-748)
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Best Local Similarity:
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AlaLeuGlyGluLeuAspIleLeuLeuGlnTrpMetGluGlu 142
                                                                                                                                    GCCACCAGAGTCATCCATGACAACTATGATCAGCTGGAGGTCCACGCTGCTGCCATTAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CAAACCCCAAAATCTTGAGAAAAATCAGCAGCATTGCCAACTCTTTCCTNTACATGCAG
                                                                                                                                                                                                                                                                                                                                                                        LysAspLeuArgLeuCysHisAlaHisMetThrCysHisCysGlyGluGluAlaMetLys 108
                                                                                                                                                                                                                  LysTyrSerGlnIleLeuSerHisPheGluLysLeuGluProGlnAlaAlaValValLys 128
                                                                                                                                                                                                                                                                                                         ProAspHisTyrThrLeuArgLysIleSerSerLeuAlaAsnSerPheLeuThrIleLys
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGCGTGACCAAGAACCTCCTGGCGTTCTACGTGGACAGGGTGTCAAAGGATCATCAGGAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AspIleArgIleLeuArgArgThrGluSerLeuGlnAspThrLysProAlaAsnArgCys
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/note="Organ: Lung; Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site_1: EcoR I; Site_2: Not I; NCI_CGAP_FT1 is a normalized cDNA library constructed from a pool of 81 RNA samples from Alveolar Macrophages
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TAG_TISSUE=Human Lung Alveolar Macrophage
TAG_LIB=UI-H-FT1
TAG_SEQ=GGCCATGCCG"
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/lab_host="DH10B (Life_Technologies)"
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/tissue_type="Alveolar Macrophage"
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Conservative:
Mismatches:
Indels:
                                                                                                                                    445
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                                                                                                                                                                                                                                                                                                                                                                                                                                  source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Laboratory for Genome Exploration Research Group, RIKEN Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-Tel: 81-45-503-9222
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Contact: Yoshihide Hayashizaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Konno,H., Fukunishi,Y., Shibata,K., Itoh,M., Carninci,P., Sugahara,Y. and Hayashizaki,Y.
Computer-based methods for the mouse full-length cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Please visit our web site (http://genome.gsc.riken.go.jp) for further details.
e mouse tissues.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Computational Analysis of Full-Length Mouse cDNAs Compared with Human Genome Sequences. Mamm. Genome. 12, 673-677 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           encyclopedia: real-time sequence clustering for construction of ononredundant cDNA library. Genome Res. 11 (2), 281-289 (2001) Kondo, S., Shinagawa, A., Saito, T., Kiyosawa, H., Yamanaka, I., Aizawa, K., Fukuda, S., Hara, A., Itoh, M., Kawai, J., Shibata, K. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RIKEN integrated sequence analysis (RISA) system--384-format sequencing pipeline with 384 multicapillary sequencer. Genome 10 (11), 1757-1771 (2000)
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BB639581.1 GI:16474967
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TCCCTGGGAGAGCTCGACGTCTTTCTAGCCTGGATTAATAAG 403
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l: genome-res@gsc.riken.jp, URL:http://genome.gsc.riken.jp/
.nci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K.,
/note="Site 1: Sall; Site_2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              709 bp mRNA linear
RIKEN full-length enriched, 7 days neona
cDNA clone A730059E15 5', mRNA sequence.
                                                                                                                                                                                       /tissue_type="cerebellum"
/dev_stage="7 days neonate"
/lab_host="DH10B"
                                                                                                                                                                clone_lib="RIKEN full-length enriched, 7 days neonate
                                                                                                                                                                                                                                                                                                    /clone="A730059E15"
                                                                                                                                                                                                                                                                                                                                                                                             organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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_xref="taxon:10090"
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RESULT 13
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Query Match:
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TITLE
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                                                   National Cancer Institute / NIH Bldg. 31 Rm10A07 Bethesda, MD 20892
                                                                                     Contact: Daniela S. Gerhard, Ph.D. Office of Cancer Genomics
                                                                                                                                                                                                                                                  Xenopus laevis (African clawed frog)
Xenopus laevis
                                                                                                                                                                                                                                                                                                      CK800046.1 GI:42812042
                                                                                                                                                                                                                                                                                                                                                          CK800046 897 bp mRNA linear EST 25-
AGENCOURT_18791077 NICHD_XGC_Te2N Xenopus laevis cDNA clone
                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Igor B. Dawid
                                                                                                                        Unpublished (1999)
                                                                                                                                     NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection
                                                                                                                                                                            Xenopodinae; Xenopus; Xenopus.
1 (bases 1 to 897)
                                                                                                                                                                                                                   Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea;
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                                                                                                                                                                                                                                                                                                                                      IMAGE:7203819 5', mRNA sequence
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cDNA Library Preparation: Express Genomics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ThrLeuArgLysIleSerSerLeuAlaAsnSerPheLeuThrIleLysLysAspLeuArg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HisLeuLeuArgLeuTyrLeuAspArgValPheLysAsnTyrGlnThrProAspHisTyr 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTGAACATACTTTTAGCCTGGATTGAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CGATGTCAGGTGCACAGACAATGTAACTGCAGTCAGGAAGCCACCAATGCAACTAGGATC 272
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                                                                                                        141 GluGlu 142
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                                                                                                                                                                                                                                        591
                                                                                                                                                                                                                                                                                                                          531 GCTAACTCTGTCCTGGGTCTAAAGATAGAGCTCAAGCATTGTCATTCTTCTATGAGGTGT
                                                                                                                                                                                                                                                                                                                                                                                                            471 AATATTTTCCCCAAAACTGGAAATCTCTTCAATTAAGGAGCAGAAAGGTCTTAATCATTTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                411 CAGATACATCCATCTGAGCAATGCTGCTTCCTTCAAACTGGGACGTTTTTACATGAAC
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                                                                                                                                                                                                                                                                                101 HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
                                                                                                                                                                                                                                                                                                                                                                  81
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      41 AspThrLysProAlaAsnArgCysCysLeuLeuArgHisLeuLeuArgLeuTyrLeuAsp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21 ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM15082 row: e column: 01
                                                                                                                                                  GAGACAGAAGCAGCTATTATTAAAGCTATTGGAGACTTGAATATCCTGATTCGCTGGTTG
                                                                                                                                                                                                                                      CCCTGTGGTGATCAGTCACACAAAATTATGGAAGACTTTAGGGAAACCTTTTACCAGATG
                                                                                                                                                                                                                                                                                                                                                             AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      was primed using oligo-dT primer:
5'-pGACTAGTTCTAGATCGCGAGCGGCCGCCC(T)25-3' and cloned into
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/lab homt="IBH10B TonA"
/clome_lib="NICHD XGC_Te2N"
/clome_lib="NICHD XGC_Te2N"
/note="Organ: testis; Vector: pExpress-1; Site_1: EcoRV;
Site_2: NotI; RNA obtained from 6 adult male testis. cDNA
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Percent Similarity:
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    US-10-789-129-26 (1-144) x CD366952 (1-658)
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Tissue Procurement: Dr. Gary W. Hunninghake, U of I
cDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA Sequencing by: Dr. M. Bento Soares, University of Iowa Clone Distribution: Distribution information can be found at http://genome.uiowa.edu/distribution/cgap.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hominidae; Homo.

1 (bases 1 to 658)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CD366952.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer: M13 FORWARD
                                                                                                                                                                                                                                                                                                                                                                                                       the conditions, other donor macrophages in different conditions). The mRNA samples were pooled for library construction. Control 0 hours; control 3 hours; control 24 hours; LPS 100 ng/ml, 3 hours; LPS 100 ng/ml, 24 hours; LPS 100 ng/ml, 3 hours; MA 10 ng/ml, 24 hours; Klebsiella moi 10, 3 hours; Klebsiella moi 10, 24 hours; Staph aureus moi 10, 3 hours; Staph aureus moi 10, 3 hours; Staph aureus moi 10, 3 hours; Staph adenoviral vector (Ad5 CMV eGFP), moi 500, 3 hours; wt adenovirus moi 500, 3 hours; wt adenovirus moi 500, 24 hours; wt adenovirus moi 500, 3 hours; wt adenovirus + LPS 3 hours; wt adenovirus + LPS 3 hours; wt adenovirus + LPS 3 hours; wt adenovirus + LPS 24 hours.

The library was subtracted according to Bonaldo, Lennon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone lib="NCI_CGAP_FT2"
/note-Torgan: Lung; Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site_1: SCOR ; Site_2: Not I;
NCI_CGAP_FT2 is a subtracted cDNA library_constructed from a pool of 81 RNA samples from Alveolar Macrophages challenged with different treatments. The mRNA samples
                                                                                                                                                                                                                                                              TAG_TISSUE=Human Lung
TAG_LIB=UI-H-FT2
TAG_SEQ=GGCCATGCCG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   were a mixture of these conditions (times refer to invage)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (some normal donor macrophages were cultured in some of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="Alveolar Macrophage"
/dev_stage="Adult"
/lab_host="DH10B (Life Technologie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="UI-H-FT2-bjp-i-22-0-UI"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Homo sapiens"
                                                    3.65e-16
217.00
68.24%
48.24%
28.82%
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                                                                                                                                                                                                                                                                                                                                                             library was subtracted according to Bonaldo, Lennon Soares, Genome Research, 6:791-806, 1996. The tissue provided by Dr. Gary W. Hunninghake of the University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (human)
                                                                     Conservative: Mismatches: Indels:
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                                                                                                                                              Length:
Matches:
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SOURCE
ORGANISM
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AUTHORS
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LOCUS
                                                                                                                                                                                                                                                                   FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Perry J. Blackshear
Office of Clinical Research and Laboratory of Signal Transduction
National Institute of Environmental Health Sciences
A2-05 NIEHS, 101 Alexander Drive, Research Triangle Park, NC 27709,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Blackshear, P.J., Lai, W.S., Thorn, J.M., Kennington, E.A., Staffa, I Jr., Moore, D.T., Bouffard, G.G., Beckstrom-Sternberg, S.M., Touchman, J.W., Bonaldo, M.F. and Bergy, M.B.
The NIEHS Kenopus maternal EST project: interim analysis of the first 13,879 ESTs from unfertilized eggs
Gene 267 (1), 71-87 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AW637118 538 bp mRNA linear bl54f02.w1 Blackshear/Soares normalized Xenopus egg laevis cDNA clone PBX0054f02 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         phone 800-533-4363 ext.cdna, fax 256-536-9016 att:cdna, email cdna@resgen.com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clone is available through Reparkway, Huntsville, AL 35901
                                                                                                                                                                                                                                                                                                                                      FORWARD: TGTAAAACGACGGCCAGT
BACKWARD: CAGGAAACAGCTATGAC
                                                                                                                                                                                                                                                                                                                                                                                   PCR PRimers
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1 (bases 1 to 538)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EST
                                                                                                                                                                                                                                                                                       Seq primer:
                                                                                                                                                                                                                                                                                                                  Plate: 0054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: black009@niehs.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fax: 919 541-4571
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Xenopus laevis
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                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA Sequencing and analyses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 919 541-4899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kenopus laevis (African clawed frog)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GluLysLeuGluProGlnAlaAlaValValLysAlaLeuGlyGluLeuAspIleLeuLeu 137
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MetThrCysHisCysGlyGluAlaMetLysLysTyrSerGlnIleLeuSerHisPhe 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SerSerLeuAlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHis
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                                                                                                                                                                                                                                                                                                                                                                                                              Intramural
    /cerl_type="unfertilized egg"
/dev_stage="unfertilized egg"
/lab_host="DH10B"
/clone_literates
                                                                                                                                                               /organism="Xenopus laevis"
/mol_type="mRNA"
/db_xref="taxon:8355"
                                                                                                                     'sex="female"
                                                                                                                                                                                                                                                           Jocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                      CAGGAAACAGCTATGACC
clone_lib="Blackshear/Soares normalized Kenopus
                                                                                                                                                clone="PBX0054F02"
                                                                                                                                                                                                                                                                                       T7 primer
                                                                                                                                                                                                                                                                                                                row: F
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library Xenopus

EST 26-APR-2001

418

538 97 598

Euteleostomi;

Pipidae;

Staffa, N.G.

Institutes

of,

Memorial

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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
Search completed: December Job time: 3796 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-789-129-26 (1-144) x AW637118 (1-538)
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                                                                                                                                                                                                               104 GluGluAlaMetLysLysTyrSerGlnIleLeuSerHispheGluLysLeuGluProGln 123
                                                                                                                                                                                                                                                                                                                                                202 CCAAAACTGGAAATCTCTTCAATTAAGGAGCAGAAAGGTCTTAATCATTTÄĞCTÄÄCTCT 261
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                                                                                                                                                                                                                                                                                   PheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCysHisCysGly 103
                                                                                                                                                                                                                                                                                                                                                                                                                                 AAGGATGTGATTACAGATGTCAGCCTCCTG---AAAGCCAAAGTGCTGAATCAGATACAT 141
                                                                                                                                                                      GATCAGTCACACÁÁAATTATGGAAGACTTTAGGGAAACCTTTTACCAGATGGAGACAGAA 381
                                                                                                                                                                                                                                                            GTCCTGGGTCTAAAGATAGAGCTCAAGCATTGTCATTCTTCTATGAGGTGTCCCTGTGGT
                                                                                                                                                                                                                                                                                                                                                                                        LysAsnTyrGlnThrProAspHisTyrThrLeuArgLysIleSerSerLeuAlaAsnSer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="vector: pT7T3-Pac; Site_1: EcoRI; Site_2: NotI; PolyA-selected mRNA was prepared from unfertilized Xenopus laevis eggs. The library was constructed in the vector pT7T3-Pac as described in Bonaldo, M.F., Lennon, G. and Soares, M.B. 'Normalization and subtraction: two approaches to facilitate gene discovery', Genome Research 6:791-806, 1996. The first strand synthesis used a NotI-dT18 primer; double stranded cDNAs were ligated to EcoRI adapters, digested with NotI, and directionally cloned into the NotI and EcoRI-digested pT7T3-Pac vector. The library contained approximately 7.2 X 10°5 recombinants, with average insert sizes of 1-1.5 kb."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5.55e-15
206.50
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32.77%
27.42%
                    28, 2005, 16:12:12
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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39
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Score:

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17

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Result
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Maximum DB
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Maximum Match 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Title:
Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OM protein - nucleic search, using frame_plus_p2n model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Command line parameters:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MODEL=frame+
                                                                                                                                                                                         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Ygapop 10.0,
Fgapop 6.0,
Delop 6.0,
    100.0
100.0
100.0
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/cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US098_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US098_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
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US-10-413-661-1

US-10-172-118-1186

US-10-342-887-186

US-10-789-129-1

US-10-789-251-1

US-10-789-968-1
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ALIGNMENTS

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Query Match:
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Best Local Similarity:
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US-10-321-163-1
                                                                                    Pred. No.:
                                                                                                                                      US-10-321-163-1
                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/341,783
PRIOR FILING DATE: 2001-12-17
NUMBER OF SEQ ID NOS: 43
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1
LENGTH: 926
                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Chandrasekher, Yasmin
APPLICANT: MCKERTAIN, PARTICIA
APPLICANT: MCKERTAIN, PARTICIA
TITLE OF INVENTION: METHOD FOR TREATING CERVICAL CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/10321163 Publication No. US20030161811A1
                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 01-44US
CURRENT APPLICATION NUMBER: US/10/321,163
CURRENT FILING DATE: 2002-12-17
                                                                                                                                                   ORGANISM: Homo sapiens FEATURE:
NAME/KEY: CDS
LOCATION: (45)...(572)
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                                                                  Matches:
                                 Conservative: Mismatches:
                   Indels:
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SOFTWARE: FastSEQ for Windows Version 3.

SEQ ID NO 1

LENGTH: 926

TYPE: DNA

ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (45)...(572)
US-10-413-661-1
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US-10-413-661-1
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Best Local Similarity:
Query Match:
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CURRENT APPLICATION NUMBER: US/10/413,661
CURRENT FILING DATE: 2003-04-15
PRIOR APPLICATION NUMBER: 09/199,586
PRIOR FILING DATE: 1998-11-25
PRIOR APPLICATION NUMBER: 60/666,597
PRIOR FILING DATE: 1997-11-26
PRIOR FILING DATE: 1997-11-26
NUMBER OF SEQ ID NOS: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application US/10413661
Publication No. US20030176657A1
GENERAL INFORMATION:
APPLICANT: Conklin, Darrell C.
APPLICANT: Haldeman, Betty A.
TITLE OF INVENTION: MAMMALIAN CYTOKINE-LIKE POLYPEPTIDE-10
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Matches:
Conservative:
Mismatches:
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NUMBER OF SEQ ID NOS: 2699

SEQ ID NO 186

LENGTH: 926

TYPE: DNA
ORGANISM: Homo sapiens
ORGANISM: Homo sapiens
PUBLICATION INFORMATION:
DATABASE ECCESSION NUMBER: AF224266
DATABASE ENTRY DATE: 2001-06-18
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APPLICANT: He, Yudong
APPLICANT: He, Yudong
APPLICANT: Linsley, Peter
APPLICANT: Mao, Mao
APPLICANT: Mao, Mao
APPLICANT: Roberts, Chris
APPLICANT: Van de Vijver, Marc
APPLICANT: Van de Vijver, Marc
APPLICANT: Bernards, Rene
TITLE OF INVENTION: Diagnosis and Prognosis of
FILE REFERENCE: 9301-175-999
CURRENT APPLICATION NUMBER: US/10/172,118
CURRENT FILING DATE: 2002-06-14
PRIOR APPLICATION NUMBER: 60/380,770
PRIOR FILING DATE: 2002-05-14
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                                                            US-10-789-129-26 (1-144) x US-10-172-118-186
                                                                                                           Query Match:
                                                                                                                            Best Local Similarity:
                                                                                                                                                               Score:
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                                                                                                                                             Percent Similarity:
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TGTGTGATCGCCACAAACCTTCAGGAAATACGAAATGGATTTTCTGACATACGGGGCAGT
                CysvalileAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer 20
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o. US20030224374A1
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Query Match:
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                                                                                                                                                                                 Score:
                                                                                                                                                                                               Alignment Scores: Pred. No.:
                                                                                                                                                                                                                                         US-10-342-887-186
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US-10-342-887-186
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LENGTH: 926
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TITLE OF INVENTION: Diagnosis and Prognosis of Breast Cancer Patients
FILE REFERENCE: 9301-188-999
CURRENT APPLICATION NUMBER: US/10/342,887
CURRENT FILING DATE: 2003-01-15
PRIOR APPLICATION NUMBER: 60/298,918
PRIOR APPLICATION NUMBER: 60/298,710
PRIOR APPLICATION NUMBER: 60/380,710
PRIOR FILING DATE: 2002-05-14
PRIOR FILING DATE: 2002-05-14
PRIOR FILING DATE: 2002-05-14
PRIOR FILING DATE: 2002-06-14
NUMBER OF SEQ ID NOS: 2699
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APPLICANT:
APPLICANT:
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APPLICANT: He, Yudong
APPLICANT: Linesley, Peter S.
APPLICANT: Mao, Mao
                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
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Roberts, Christopher J.
Van 't Veer, Laura Johanna
Van de Vijver, Marc J.
Bernards, Rene
                                         CysValIleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer 20
ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln 40
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Query Match:
DB:
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                                                                                                              US-10-789-129-26 (1-144) x US-10-789-129-1 (1-926)
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Publication No. US20040142428A1
GENERAL INFORMATION:
APPLICANT: Conklin, Darrell C.
APPLICANT: Haldeman, Betty A.
TITLE OF INVENTION: MAMMALIAN CYTOKINE-LIKE POLYPEPTIDE-10
FILE REFERENCE: 97-72
CURRENT APPLICATION NUMBER: US/10/789,129
CURRENT FILING DATE: 2004-02-27
                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 43
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 1
LENGTH: 926
                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 09/199,586
PRIOR FILING DATE: 1998-11-25
PRIOR APPLICATION NUMBER: 60/066,597
PRIOR FILING DATE: 1997-11-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION NUMBER: US/10/413,661
PRIOR FILING DATE: 2003-04-15
                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
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               ValGlnAlaLysAspGlyAsnIleAspIleArgIleLeuArgArgThrGluSerLeuGln 40
                                                                    CysVallleAlaThrAsnLeuGlnGluIleArgAsnGlyPheSerAspIleArgGlySer
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APPLICANT: CONKIIO.

APPLICANT: CONKIIO, Darrell C.

APPLICANT: Haldeman, Betty A.

TITLE OF INVENTION: MAMMALIAN CYTOKINE-LIKE POI

FILE REFERENCE: 97-72

CURRENT APPLICATION NUMBER: US/10/789,251

CURRENT FILING DATE: 2004-02-27

PRIOR APPLICATION NUMBER: US/10/413,661

PRIOR FILING DATE: 2003-04-15

PRIOR FILING DATE: 1998-11-25

PRIOR APPLICATION NUMBER: 09/199,586

PRIOR FILING DATE: 1998-11-25

PRIOR FILING DATE: 1998-11-26

PRIOR FILING DATE: 1997-11-26

NUMBER OF SEQ ID NOS: 43

SOPTWARE: FASTSEQ for Windows Version 3.0

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LENGTH: 926
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US-10-789-251-1
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Publication No. US20040143097A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORCANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (45)...(572)
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                   GACACAAAGCCTGCGAATCGATGCTGCCTCCTGCGCCATTTGCTAAGACTCTATCTGGAC 320
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FILE REFERENCE: 97-72
CURRENT APPLICATION NUMBER: US/10/789,968
CURRENT FILING DATE: 2004-02-27
PRIOR APPLICATION NUMBER: US/10/413,661
PRIOR APPLICATION NUMBER: 09/199,586
PRIOR APPLICATION NUMBER: 09/199,586
PRIOR APPLICATION NUMBER: 60/066,597
PRIOR APPLICATION NUMBER: 60/066,597
PRIOR PILING DATE: 1997-11-26
NUMBER OF SEQ ID NOS: 43
SOFTWARE: FASUSEQ for Windows Version 3.0
SEQ ID NO 1
LENGTH: 926
TYPE: DNA
ORGANISM: Homo sapiens
PEATURE:
NAME / NEW. COS
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Conklin, Darrell C.
APPLICANT: Haldeman, Betty A.
TITLE OF INVENTION: MAMMALIAN CYTOKINE-LIKE POLYPEPTIDE-10
                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: CDS
LOCATION: (45)...(572)
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Sequence 1, Application US/10748484

Publication No. US20040181040A1

GENERAL INFORMATION:
APPLICANT: Conkin, Darrell C.
APPLICANT: Haldeman, Betty A.
APPLICANT: Haldeman, Angelika
TITLE OF INVENTION: MAMMALIAN CYTOKINE-LIKE POLYPEPTIDE-10

FILE REFERENCE: 97-72

CURRENT APPLICATION NUMBER: US/10/748,484

CURRENT FILING DATE: 2003-12-29

PRIOR APPLICATION NUMBER: US/09/199,586

PRIOR FILING DATE: 1998-11-26

NUMBER OF SEQ ID NOS: 43

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 1

LENGTH: 926

TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (45)...(572)

US-10-748-484-1
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US-10-748-484-1
Sequence 1, Ap
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APPLICANT: CA
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CATTOTON NUMBER: US 60/524.131 GANTE 200-21.2 GASSAGO NUMBER: US 60/524.131 GANTE 200-31.2 GATTON NUMBER: US 60/524.131 GANTE 200-31.2 GATTON NUMBER: US 60/524.131 GANTE 200-31.2 GATTON Sapiens 8. 29e-93 Length: 100.00% Illarity: 100.00% Ill	AGAAGCAATGAAGAATACAGCCAGATT AGGAAGCAATGAAGAAATACAGCCAGATT AGGAAGCAATGAAGAAATACAGCCAGATT AGGAAGTTGTGAAGGCTTTGGGGGAACT US/10994116 1136004A1 A 572 Wayne even D. her, Yasmin A. acey R. yce M. windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0 Windows Version 4.0	1 1 9 ma o 6 m P 9 1 FERRITA NATIONAL SERVICE	121 G1 501 GA 601
Ser HGT	994-116-1 (1-926) euGlnGluIleArgAsnG TCAGGAAATACGAAATG	1 CysVallleAlaThrAsnLe	-10-789-
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	/10/994,116 19 0/524,131 0/555,857 Version 4.0	PERRENCE: 03-21 APPLICATION NUMBER: US, FILING DATE: 2004-11-; PPLICATION NUMBER: US 6; PPLICATION NUMBER: US 6; PPLICATION NUMBER: US 6; PPLICATION NUMBER: US 6; ILING DATE: 2004-03-24 DF SEQ ID NOS: 90 D1 1905-1905-1906-1906-1906-1906-1906-1906-1906-1906	FILE RE CURRENT CURRENT CURRENT PRIOR A PRIOR F PRIOR F PRIOR F NUMBER (SOFTWARK
BINDING PARTNERS AND METHODS OF	AND	NT: Xu, Wenfeng NT: Kindsvogel, Wayne NT: Chen, Zhi NT: Hughes, Steven D. NT: Hughes, Stacey R. NT: Dillon, Stacey R. NT: Lehner, Joyce M. NT: Siadak, Anthony W. NT: Sivakumar, Pallavu NT: Moore, Margaret D. F INVENTION: ANTI-IL-20 F INVENTION: INFLAMMAT.	APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA
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Sequence 1, Application US/10994151

Publication No. US20050170468A1

GENERAL INFORMATION:
APPLICANT: Ku, Wenteng
APPLICANT: Chen, Zhi
APPLICANT: Chen, Zhi
APPLICANT: Chen, Zhi
APPLICANT: Chen, Zhi
APPLICANT: Chandrasekher, Yasmin A.
APPLICANT: Lehner, Joyce M.
APPLICANT: Lehner, Joyce M.
APPLICANT: Siadak, Anthony W.
APPLICANT: Sivakumar, Pallavur V.
APPLICANT: Moore, Margaret D.
TITLE OF INVENTION: ANTI-IL-20 RECEPTOR ANTIBODIES AND BINDING PARTNERS AND METHODS C
TITLE OF INVENTION: IN INFLAMMATION
FILE REPERENCE: 04-17
CURRENT APPLICATION NUMBER: US/10/994,151
CURRENT FILING DATE: 2004-11-19
DEFIOR ADDITION TO MIMBER: US/10/994,151
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US-10-994-151-1
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 926
TYPE: DNA
ORGANISM: Homo
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PRIOR FILING DATE: 2003-11-21
PRIOR APPLICATION NUMBER: US 60/555,857
PRIOR FILING DATE: 2004-03-24
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LOCATION: (45)...
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APPLICANT: CHAN, CHUNG

TITLE OF INVENTION: METHODS FOR ENHANCING THE TRANSLATION

TITLE OF INVENTION: AND EXPRESSION OF RECOMBINANT PROTEINS

FILE REFERENCE: 01-15

CURRENT APPLICATION NUMBER: US/10/122,882B

CURRENT APPLICATION NUMBER: 00/283,668

PRIOR APPLICATION NUMBER: 60/283,668

PRIOR APPLICATION NUMBER: 60/283,668

PRIOR FILING DATE: 2001-04-13

NUMBER OF SEG ID NOS: 25

SOFTWARE: FASTSEQ FOR WINDOWS Version 4.0

SEG ID NO 14

LENGTH: 462

TYPE: DNA

ORGANISM: Homo sapiens

FEATURE:

NAME YOUNG THE TRANSLATION

PRATURE:

NAME YOUNG THE TRANSLATION

ORGANISM: Homo sapiens
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Best Local Similarity:
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CURRENT FILING DATE: 2005-02-09
PRIOR APPLICATION NUMBER: 11/008,853
PRIOR FILING DATE: 2004-12-10
PRIOR APPLICATION NUMBER: US 60/529,412
PRIOR FILING DATE: 2003-12-12
NUMBER OF SEQ ID NOS: 29
SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Pownder, Tracey
TITLE OF INVENTION: METHODS FOR ENHANCING
TITLE OF INVENTION: RECOMBINANT PROTEINS
FILE REFERENCE: 03-22C1
                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens
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GluGluThrGlu
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APPLICANT: POWNDER, TRACEY
APPLICANT: CHAN, CHUNG
TITLE OF INVENTION: METHODS FOR ENHANCING THE TRANSLATION
TITLE OF INVENTION: AND EXPRESSION OF RECOMBINANT PROTEINS
FILE REFERENCE: 01-15
FULRENT APPLICATION NUMBER: US/10/122,882B
CURRENT FILING DATE: 2002-04-12
PRIOR APPLICATION NUMBER: 60/283,668
PRIOR FILING DATE: 2001-04-13
NUMBER OF SEQ ID NOS: 25
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 17
LENGTH: 513
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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  GAACCGCAGGCAGCAGTGGTGAAAGCTCTGGGTGAACTGGACATTCTGCTGCAGTGGATG
                                                                               CACTGTGGTGAGGAAGCAATGAAAAAATACAGCCAGATTCTGAGCCACTTCGAAAAACTG
                                                                                                         HisCysGlyGluGluAlaMetLysLysTyrSerGlnIleLeuSerHisPheGluLysLeu
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                                                                                                                                                                                    AlaAsnSerPheLeuThrIleLysLysAspLeuArgLeuCysHisAlaHisMetThrCys
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GAGGAGACCGAA 510

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Sequence 1. Application US/10467431

Publication No. US20040092445A1

GENERAL INFORMATION:

APPLICANT: Liu, Ling

APPLICANT: Liu, Ling

APPLICANT: Timothy, Noblitt

ITILE OF INVENTION: USE OF LP82 TO TREAT HEMATOPOIETIC DISORDERS

FILE REFERENCE: X-14753

CURRENT FILING DATE: 2003-08-06

PRIOR APPLICATION NUMBER: US/10/467,431

CURRENT FILING DATE: 2001-02-28

PRIOR APPLICATION NUMBER: 60/272,242

PRIOR APPLICATION NUMBER: 60/272,740

PRIOR APPLICATION NUMBER: 60/332,740

PRIOR FILING DATE: 2001-11-19

NUMBER OF SEG ID NOS: 5

SOFTWARE: PatentIn version 3.1

LENGTH. 667
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TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (272)...(799)
OTHER INFORMATION:
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NAME/KEY: sig_peptide
LOCATION: (272)..(325)
OTHER INFORMATION:
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                     GluGluThrGlu 144
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PRIOR APPLICATION NUMBER: 60/264,239
PRIOR FILING DATE: 2001-01-26
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PatentIn version 3.2
SEQ ID NO 1
LENGTH: 867
TYPE: DNA
ORGANISM: Homo sapiens
PEATURE:
NAME/KEY: CDS
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Publication No. US20040162237A1
GENERAL INFORMATION:
APPLICANT: FOX, NYIES W
TITLE OF INVENTION: USE OF LP82 TO TREAT BODY WEIGHT DISORDERS
FILE REFERENCE: X-14544
CURRENT APPLICATION NUMBER: US/10/466,695A
CURRENT FILING DATE: 2004-02-03
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NAME/KEY: sig_peptide
LOCATION: (272)..(325)
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